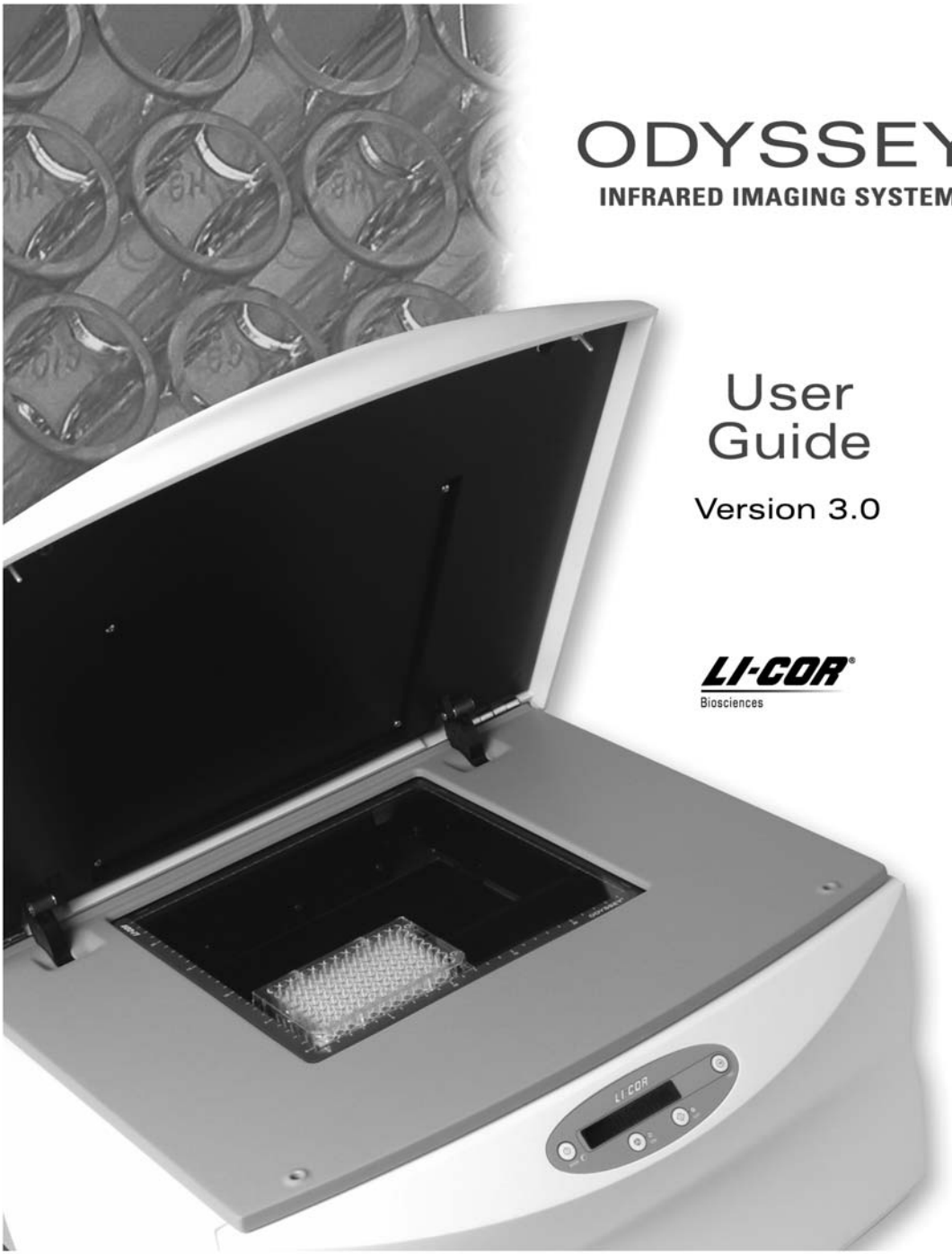
ODYSSEY

INFRARED IMAGING SYSTEM

User Guide

Version 3.0

LI-COR Biosciences



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7.3. If these representatives have not agreed on a resolution of such dispute within ten (10) Business Days of its referral to them, the dispute shall be promptly submitted to a neutral adviser (the "Adviser") who shall be chosen from the list of arbitrators registered with the American Arbitration Association. For purposes of this Section, "Business Day" shall mean each weekday and the hours of such weekday in

which Licensee is open for business. The Adviser shall, within fourteen (14) days of the submission, recommend, in writing, a procedure for resolving the dispute and shall specify in such writing whether such procedure shall be binding, non-binding or involve a combination of binding and non-binding procedures.

7.4. If the parties do not mutually agree upon the process recommended b the Adviser within ten (10) Business Days of their receipt of the Adviser's written recommendation, they shall promptly convene a non-binding hearing (the "Mediation"). The rules for Mediation will be established by the Adviser, after consultation with the parties.

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Chapter 1: Introduction

How to Learn Odyssey

If you are upgrading from a previous version of Odyssey® software, a list of changes for version 3.0 can be found in the help system. In addition, movies that illustrate new software features can be viewed by choosing **Help > What's New**.

The best way for new users to learn Odyssey is to work through the tutorials in the **Tutorial Manual**. The Tutorial Manual is a step-by-step guide that introduces you to scanning with the Odyssey Imager, as well as analysis with Odyssey software. The overview of Odyssey in the Tutorial Manual will familiarize you with basic operation of the entire system.

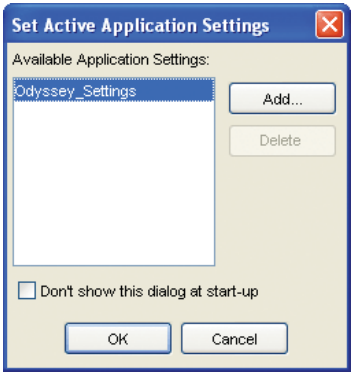
When you are ready for more information, this **User Guide** is a reference manual with complete descriptions of sizing and quantification, as well as features of the Odyssey In-Cell Western Module. The **Odyssey In vivo Imaging Guide** describes optional software module and operational details for scanning mice using the Odyssey MousePOD™ Imaging Accessory.

Sample preparation is described in the Odyssey **Application Protocols Manual** and in the pack inserts enclosed with reagents.

Operation and maintenance of the Odyssey instrument can be found in the Odyssey **Operator's Manual**. Documentation of the server software inside the Odyssey instrument is also included in the Operator's manual. User account management, networking, troubleshooting, scan control, and software updates are all discussed.

Selecting an Application Settings File at Start-up

When the Odyssey application starts, a window is displayed that asks the user to choose the application settings file for the current session. This makes it easy for users to have their own settings file that determines important parameters such as background calculation method. Until you understand the Odyssey application, it is best to choose the default application settings stored in **Odyssey_Settings** as shown below. The active settings file can be changed at any time by choosing **Settings > Select Active Settings**. Chapter 12 describes adding, deleting and changing the active application settings file.

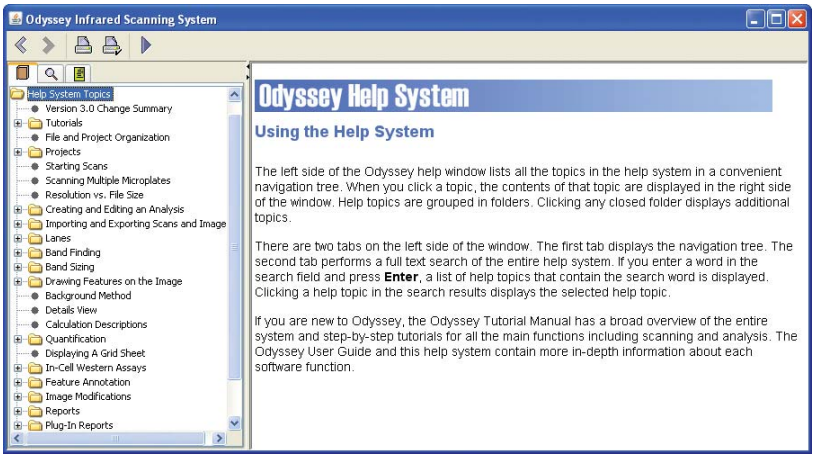


Setting Up Users and Scanners

Before you can do anything in Odyssey software, you must have your own user account. The **Settings** menu is used to to add user accounts. Chapter 12 describes this procedure. The Odyssey instrument network address must also be added (**Settings > Scanners**) to Odyssey software before the instrument can be operated. Odyssey instruments (scanners) are generally added during installation, but Chapter 12 describes this function should you need to perform it.

The Odyssey Help System

The Odyssey help system can be invoked by choosing **Help > Contents** or by pressing **F1** on the keyboard.



The Help window has two frames. The left frame contains navigational links. Click a topic to display content for the topic in the right frame. Folders in the navigational frame contain additional topics and are opened by clicking their *plus* symbol. To search for something specific, click the search tab (magnifying glass icon) and enter the search text.

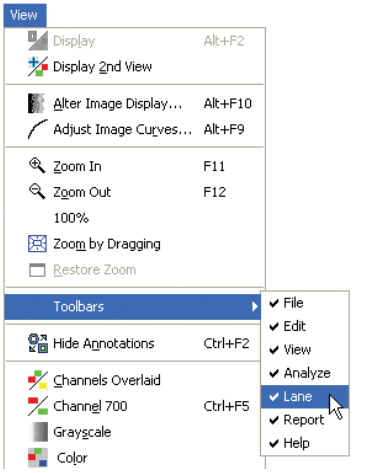
The help system contains most of the information found in the Tutorial Manual and this User Guide. However, the information is organized in a more task-oriented way that should help if you forget how to do something.

Toolbars

The Odyssey user interface makes extensive use of toolbars to provide single-click access to most functions. The function of each tool is given in a tool tip that can be displayed by stopping the cursor over the tool on the toolbar. A description of each tool can also be found in the online help system.

Most tools on the toolbar corresponds to a menu choice on the menu bar that does the same function. The examples throughout this manual use both the toolbar and menu functions.

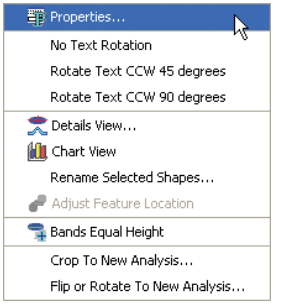
If the toolbars get in your way, you can easily hide them. To hide a toolbar, choose **View > Toolbars** and deselect the toolbar you want to hide.



Context-Sensitive Menus

Odyssey has context-sensitive menus that change depending on what is selected when the menu is opened. To open a context-sensitive menu select a feature on the image, such as a band marker, and right-click the image.

Using context sensitive menus you can do things like open the Properties window for an object, rotate text annotations, and plot a histogram of quantification values.



Correcting Mistakes

Odyssey software has extensive "Undo" capabilities that are accessed on the **Edit** menu. By continuing to choose **Edit > Undo** or clicking the tool, you can undo the last 100 operations since the Odyssey program was opened (with a few exceptions). The number of undo's can be changed in the Application Settings (choose **Settings > Application** and select **General** from the settings list).

Odyssey Project and File Organization

Projects, Scans, and Analyses

Odyssey file organization starts with the *project* folder. A *project* folder is a folder anywhere on a local or network drive that is used to store Odyssey scans. *Project* folders can be used to separate scans into a logical structure that compliments your research. *Project* folders are created when new projects are started (choose **File > New**).

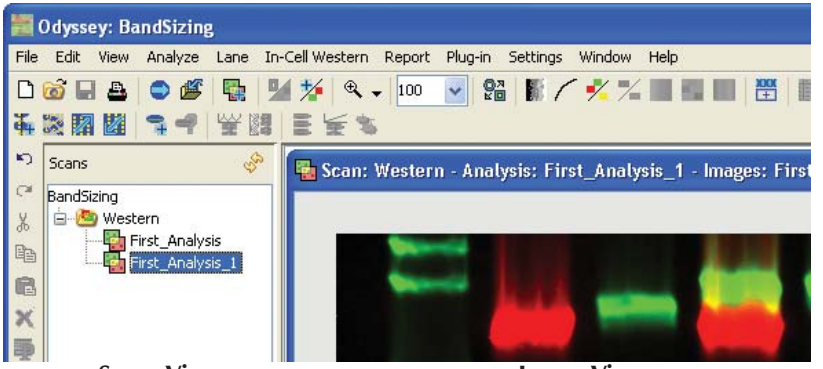
Within a project folder there can be many *scans*. Each *scan* is a folder containing one or two TIFF images from the Odyssey Imager, depending on whether probes for one or both dyes were used. *Scans*

can be started from the New Project window or by clicking the Scan button on the toolbar ( ) if a project is already open.

Scan folders also contain the *analysis* files generated when an analysis is performed on the scan. *Analysis* files hold all the data (concentrations, etc.) and annotations created when the scan was analyzed. At the end of each new scan, the first *analysis* on the new image files is saved. A set of images can be analyzed as many times as needed. A new analysis can be created by clicking the New Analysis button on the toolbar ( ).

Displaying Projects, Scans, and Analyses in the Main Odyssey Window

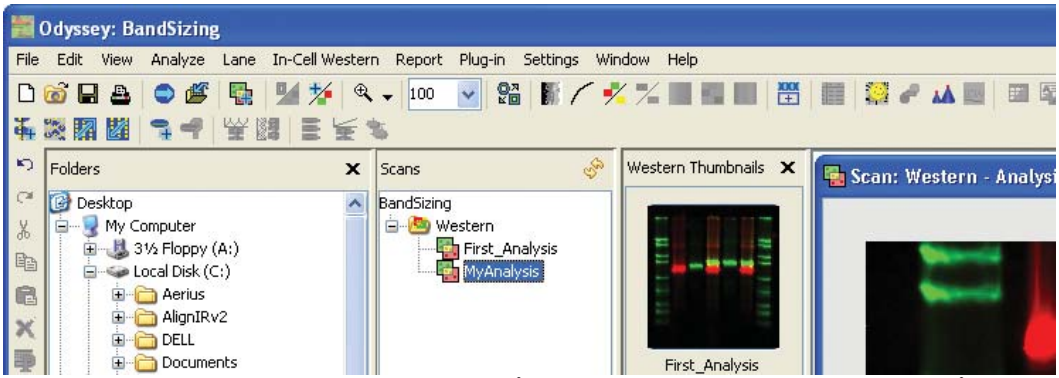
The main Odyssey window shown below is the default window configuration that displays the Scans view (left) and the Image view window (right).



Scans View **Image View**

The Scans view is always open (left) and lists the projects, scans, and analyses for the current project. Double clicking on an analysis opens an Image view (right) containing the images from the analysis.

The **View** menu can be used to display other file information in the main Odyssey window. The Folders view (shown below) displays a directory tree similar to that found in Windows Explorer. The purpose of the Folders view is to aid in finding and opening Projects, which are displayed in the Scans view (center). The Thumbnails view shows a thumbnail sized image of the scan or analysis selected in the Scans view. The Folders, Scans, and Thumbnails views are discussed below.



*Folders view and Thumbnails view can be opened using the **View** menu.*

Folders View

Folders view is opened (or hidden) by choosing **View > Folders View**. A close button in the upper right corner hides the view. Odyssey project folders, which contain scans, have a unique icon ( ), as do the scan folders ( ) within project folders. Projects that are open and have been edited are shown with a pencil icon ( ).

All folders can be expanded by clicking the "plus" icon next to the folder ( ). Folders can also be expanded by double-clicking them. When an Odyssey project folder is double-clicked, the project is opened and shown in the Scans view. The context menu that opens

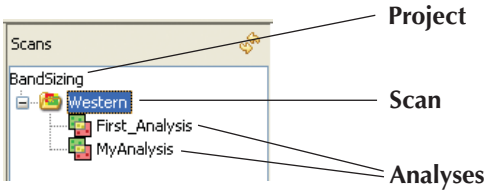
by right-clicking on a folder can also be used to open and close a folder.

Note: *Starting with Odyssey software version 3.0, project folders can be stored in any location. Local drives and mapped network drives generally offer the best performance.*

In addition to browsing for projects and scans in Folders view, a search function (**File > Scan > Search for Scans**) is available to find scans if the file name is known or partially known. The list of recently open projects on the **File** menu is also a fast way to open projects.

Scans View

The Scans view shows all scans and analyses for the current project. The name of the current project is shown in Scans view and highlighted in Folders view.



In the project named “Bandsizing” above, there is one scan named “Western” and two analyses named “First_Analysis” and “MyAnalysis”. The project above has only one scan, but for projects with many scans it may be useful to sort the scans. Right-clicking the project name opens a context menu with choices for sorting scans by name or date, using either ascending or descending order. The default sort order can be set by choosing **Settings > Application** and selecting **General** from the **Settings List**.

To view all the analyses associated with a scan, click the “plus” icon () to expand the scan folder. To open an analysis, double-click it in the scans list. The first analysis in a scan folder can be opened

by double-clicking the scan folder. An analysis can also be opened using Thumbnails view as described below.

A scan or analysis can be deleted by selecting it in the Scans view and pressing **Delete** on the keyboard. An analysis can also be deleted by right-clicking the analysis and choosing **Delete Scan** from the popup menu. Additionally, the operating system can be used to delete files in the normal way. If Odyssey software is open when files are deleted using the operating system, the Scans view will not immediately show the files have been deleted. The refresh button () in the upper right corner of Scans view can be used to refresh the scan list and show any changes.

Thumbnails View

Thumbnails view is opened (or hidden) by choosing **View > Thumbnails View**. A close button in the upper right corner hides the view. One composite, two-channel thumbnail image is shown for each analysis in the scan folder. Double-clicking a thumbnail image opens the corresponding analysis.

Initially if a scan or analysis is not selected in the scan list, the message “No Thumbnail Defined” is displayed. All the thumbnails are created as soon as a scan or analysis is selected. The thumbnails are real files that are saved as JPEG files in the scan folder. The file name convention is *ScanName_AnalysisName_tn.jpg*. If changes are made to the image, such as cropping or rotating, these changes will not be updated in the thumbnail until the analysis is saved.

Chapter 2: Starting Scans

How to Start Scans

Scans on the Odyssey Imager can be started using the Windows®-based Odyssey Software, an Internet browser, or from the front panel of the Odyssey Imager. Chapter 6 of the Odyssey Operator's Manual discusses starting scans using an Internet browser. Front panel operation is described in Chapter 7 of the Odyssey Tutorial Manual and Chapter 3 of the Odyssey Operator's Manual. The remainder of this chapter is dedicated to starting both standard scans and multiple microplate scans with Odyssey software. Scanning mice with the MousePOD™ Accessory is discussed in the Odyssey *In vivo* Imaging Guide included with the MousePOD.

Before a scan can be started, a project must be open so the new scan can be stored in the open project.

Starting Standard Scans in an Existing Project

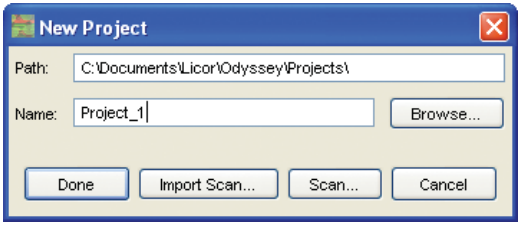
Existing projects are opened by choosing **File > Open** or by clicking  on the toolbar. The four most recently opened projects are also listed toward the bottom of the **File** menu. The number of recent projects listed can be increased to as many as 10 in the Application settings (choose **Settings > Application** and select **General** from the **Settings List**).

Once a project is open, a standard scan can be started by clicking  on the toolbar or choosing **File > Scan > Scan**. After entering

your user name and password, the Scanner Console window is opened, allowing scans to be started as described below.

Starting a Standard Scan in a New Project

To start a new project in Odyssey Software, choose **File > New**.

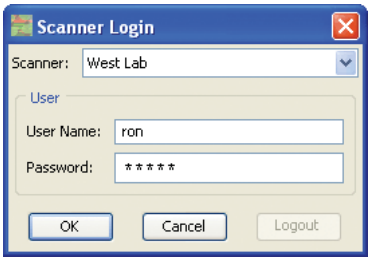


The path and project name can be entered by clicking **Browse** to open a standard "new file" window. File paths and names can also be typed in the **Path** and **Name** fields.

After entering the project name, take one of the following actions:

- Click **Done** to create an empty project.
- Click **Import Scan** to create the project and import images from a different project (Chapter 3).
- Click **Scan** to create the project and start a standard scan that will become part of the project. In the Scanner Login window, select the scanner (if necessary), enter your **User Name** and **Password**, and click **OK**.

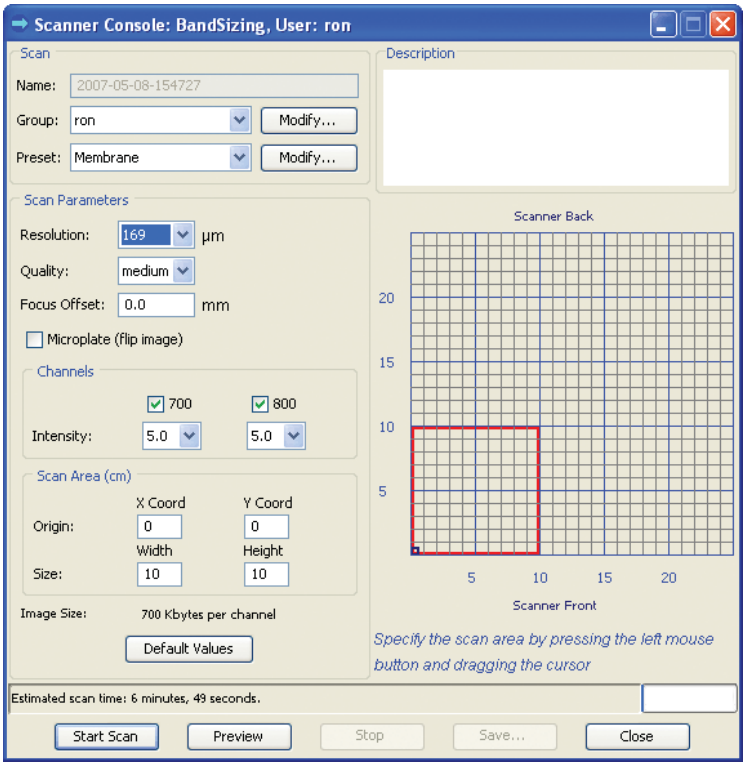
*If someone else is already logged in, click **Logout** before entering your **User Name** and **Password**.*



Note: *User names and passwords must be added by a user with Administrator (admin) access privileges. The system administration functions (**Settings > System Administration**) are used to add users and set access privileges (see Chapter 12).*

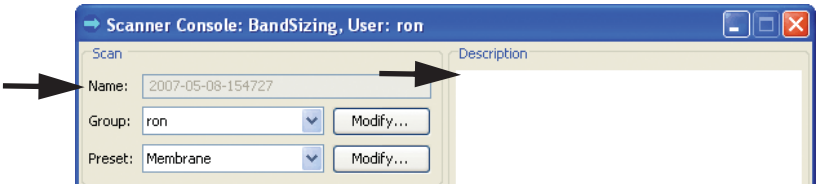
Scanner Console Window for Standard Scans

Whether a scan is started in a new or existing project, the Scanner Console window is used to specify the scan parameters and start the scan. It can also be used for a quick preview scan. During each scan, the Scanner Console displays the scan in real time as it is collected and displays progress indicators for the scan.



Naming a Scan and Entering a Description

The **Name** field is not editable at the beginning of a scan. The default scan name is filled in automatically according to the naming conventions in the Application Settings. The default name may be blank, a sequential name, or a time stamp (shown below). At the end of the scan, the default name can be accepted or replaced with a different name before the file is stored on the computer.



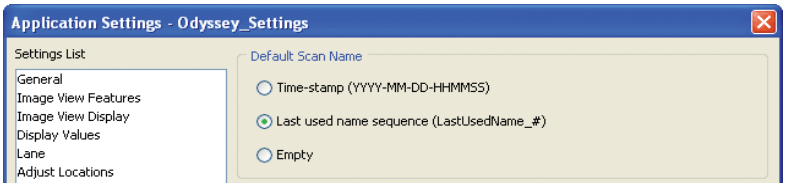
The name in the **Name** field is also the scan name that will be stored on the hard drive of the Odyssey instrument. If "blank" is the current naming convention, a time stamp will be used for the scan name on the Odyssey instrument. If the default name is replaced with a new scan name at the end of the scan, the scan name on the computer will be different than the original scan name on the hard drive in the Odyssey instrument. The original scan name can be viewed by choosing **Edit > Scan Description** to view the scan description. The original name is also listed in the tool tip that is displayed when the cursor is stopped over a scan name in the Scans view of the main Odyssey window.

Entering a description in the **Description** field is optional; however, descriptions can be included in reports.

Changing the Default Scan Name

To change the default scan name, choose **Settings > Application** and select **Naming Conventions** from the **Settings List**.

Since these are Application settings, naming conventions can be saved in the settings files for individual users (Chapter 12), allowing each user to have scan names automatically entered as desired.



Default Scan Names:

- **Time Stamp:** When the Scanner Control window is opened, the current year, month, day, hour, minutes, and seconds is entered in the **Name** field automatically. This time stamp can be edited or appended with other text.
- **Last Used Name Sequence:** When the Scanner Control window is opened, the name of the last scan is entered in the **Name** field and appended with a sequential number. For example, if MyScan was the last scan name, Odyssey will present MyScan_1 as the default name for the next scan, followed by MyScan_2, etc.
- **Empty:** When the Scanner Control window is opened, the **Name** field is left blank so the user can enter a name at the end of a scan.

Previewing a Scan (Optional)

Click **Preview** (optional) in the Scanner Console window to scan a low resolution preview before starting high resolution scanning.

A preview scan is a low resolution scan at the lowest quality setting that takes only a few minutes to complete, depending on scan area. A preview scan can be used to check fluorescent signal intensity or to adjust the scan area before high resolution scanning. Adjusting the scan area (see *Setting Scanner Parameters for Standard Scans* below) can shorten scan times by reducing the amount of empty background that is scanned.

Selecting a Scan Group

A scan group is a special directory on the Odyssey instrument that has restricted access. Initially, users have access to the *Public* scan group and a scan group that matches their user name. Additional scan groups can be created for special purposes. For example, if several people are doing scans for a particular research project, it might be useful to keep all scans for that project in one scan group.

The **Group** drop-down list is used to select the scan group in which the new scan will be stored.

Scan groups are added and deleted by clicking **Modify** (next to **Group**). See Chapter 12 for complete information on scan groups.

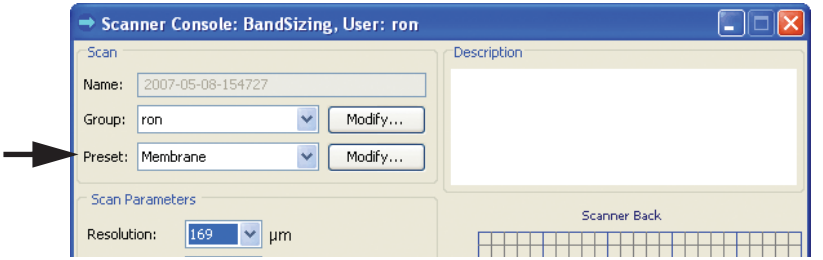
Setting Scanner Parameters for Standard Scans

Scan parameters, such as resolution and scan area, can all be entered individually in the Scanner Console window or loaded from stored sets called *Presets*. For most scans it is easiest to load Preset parameters and then edit individual parameters, such as scan area, to match the current scan.

Loading Preset Parameters

Sets of scan parameters can be chosen from the **Preset** drop-down list. When a Preset is chosen, all existing scan parameters in the

Scanner Console window are replaced by scan parameters stored in the Preset file.



Odyssey software initially has four Preset files for general use – one for membranes, two for gels, and one for microplates. In addition, there are four Preset files for the MousePod™ In Vivo Imaging Accessory: one for each of the three mouse positions in the MousePod and one for all three positions at once.

Note: *For older instruments, Odyssey Server Software version 2.0 or above is required in order to have enough focus offset to scan a microplate. See Odyssey Operator's Manual to determine software version number.*

Membrane, Gel, and Microplate Presets

	Membrane	DNA Gel	Protein Gel	Microplate2
Resolution	169	169	169	169
Quality	medium	medium	medium	medium
Focus Offset	0.0	2.0	0.5	3.0 mm
Channels	700, 800	700, 800	700, 800	700, 800
Intensity	5.0	8.0	5.0	5.0
Scan Origin	0,0	0,0	0,0	0,0
Scan Size	10,10	10,10	10,10	13,9

Note: *There are Presets both in the Odyssey Imager itself and in Odyssey Software. The Presets in the Odyssey Imager are used when starting scans from the front panel or from an Internet browser. Presets in Odyssey Software are used only in Odyssey Software. Information on using, modifying, and saving Presets in the Odyssey Imager can be found in the Odyssey Operator's Manual.*

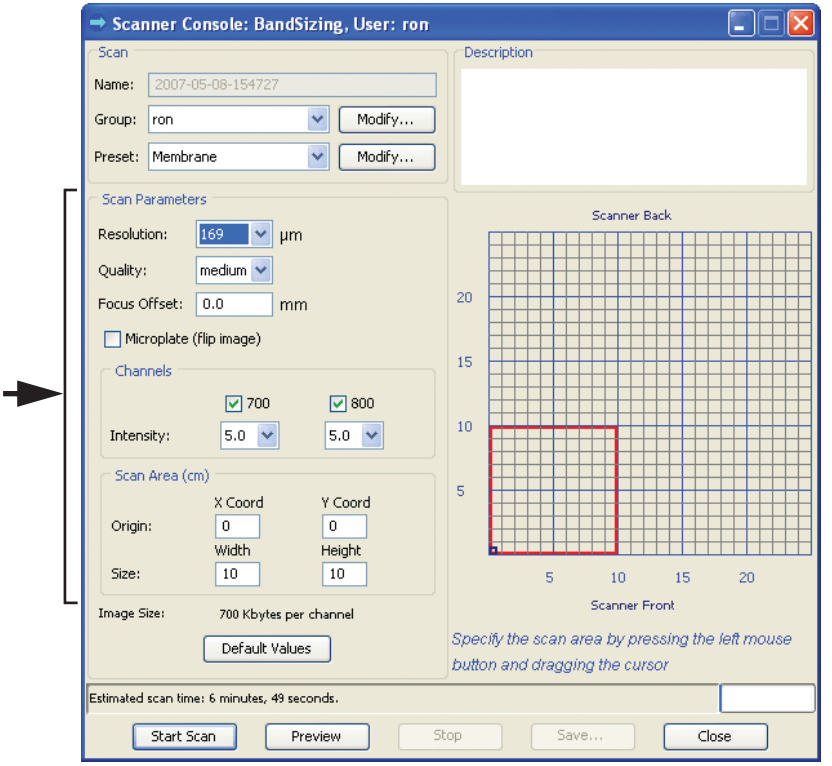
MousePOD™ Presets

	Full Pod Scan	Mouse Center Position	Mouse Left Position	Mouse Right Position
Resolution	169	169	169	169
Quality	medium	medium	medium	medium
Focus Offset	1.0 mm	1.0 mm	1.0 mm	1.0 mm
Channels	700, 800	700, 800	700, 800	700, 800
Intensity	L1.0, L2.0	L1.0, L2.0	L1.0, L2.0	L1.0, L2.0
Scan Origin	0,0	8,0	0,0	15,0
Scan Size	25,19	9,19	10,19	10,19

See Odyssey *In vivo* Imaging Guide for details on scanning with the Odyssey MousePOD Accessory.

Editing Scan Parameters for Standard Scans

All scan parameters are listed in the middle of the Scanner Console window. Each parameter can be edited as described below.



Resolution can be set to 21, 42, 84, 169, or 337 µm. For typical scans of membranes or gels, 169 µm scans should suffice. As resolution increases, file sizes get very large. The table below shows the resolution and scan size limits for starting scans with Odyssey Software.

File sizes under 7 MB per image scan well and can be analyzed in Odyssey Software without cropping the image into smaller pieces. File sizes from 7 - 14 MB are marginal and Odyssey Software may

run out of memory during a scan. Scans with file sizes larger than 14 MB per image can be performed using the browser interface as described in the Odyssey Operator's Manual, but should not be attempted with Odyssey Software. The table below shows typical combinations of resolution and scan size, with shading to indicate file sizes that are too large for Odyssey Software.

	Resolution				
Scan Size	**337 µm**	**169 µm**	**84 µm**	**42 µm**	**21 µm**
5 x 5 cm	44k	175k	708k	2.8M	11.3M
5 x 10 cm	88k	350k	1.4M	5.7M	22.6M
5 x 15 cm	132k	525k	2.1M	8.5M	34.0M
5 x 20 cm	176k	700k	2.8M	11.3M	45.3M
5 x 25 cm	220k	875k	3.5M	14.2M	56.7M
10 x 10 cm	176k	700k	2.8M	11.3M	45.3M
10 x 15 cm	264k	1.0M	4.1M	17.0M	68.0M
10 x 20 cm	352k	1.4M	5.6M	22.7M	90.7M
10 x 25 cm	440k	1.7M	7.0M	28.3M	113.3M
15 x 15 cm	396k	1.6M	6.3M	25.5M	102.0M
15 x 20 cm	528k	2.1M	8.4M	34.0M	136.0M
15 x 25 cm	660k	2.6M	10.6M	42.5M	170.0M
20 x 20 cm	704k	2.8M	11.3M	45.4M	181.4M
20 x 25 cm	800k	3.5M	14.1M	56.7M	226.7M
25 x 25 cm	1.1M	4.4M	17.6M	70.9M	283.4M

File size is small enough to scan with Odyssey Software.
Marginal for Odyssey Software.
Scan should be started in using the browser interface.

Odyssey Software also has limitations on the size of images that can be analyzed. The total size of all open images should not exceed 20-25 MB. One analysis with two 10MB images will use up most of the memory resources. However, if your typical image size is 2 MB, five separate analyses can be opened. Large scans can be cropped into smaller pieces using the browser software if necessary.

For band sizing applications, the resolution setting can be checked by looking at the lane profiles (Chapter 5). If the lane profile shows many small jagged peaks on the larger peaks of bands (as contrasted with smooth peaks), this may indicate the resolution is too coarse. These jagged peaks will influence the accuracy of band finding. If the small peaks are caused by lack of resolution, choosing a smaller resolution value should improve the problem.

Quality controls scan speed and ultimately how many detector readings are processed for a given area on the membrane in order to make one pixel on the image. For typical scans, **Medium** is recommended, but there are five settings. Choosing **Highest** quality will reduce noise in the image data, but significantly increase scanning time due to the slower scanning speed. Similarly, choosing **Lowest** will decrease scan time, but increase noise in the image data. For high resolution scans where samples have very little fluorescence, **High** or **Highest** may be a better choice than **Medium**. When **Quality** is set too low, the image may become noisy or "grainy", particularly in the background.

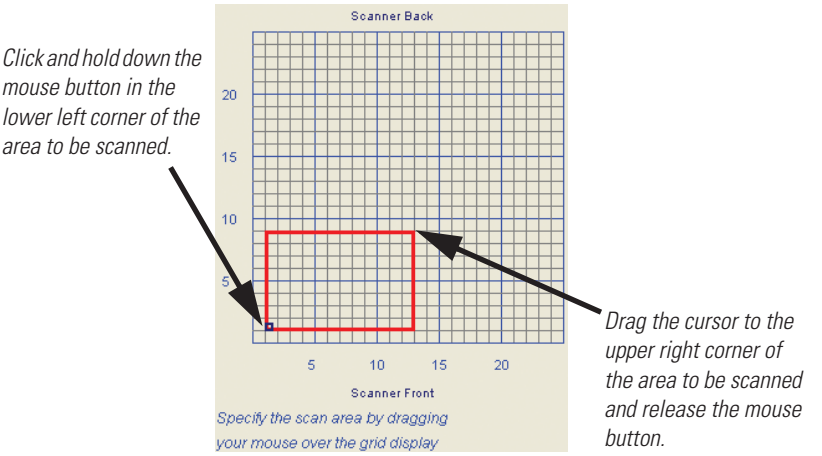
Focus Offset should always be zero when scanning membranes. For gels, set **Focus Offset** to half the gel thickness, in millimeters. For microplates recommended by LI-COR (Operator's Manual, Chapter 3), focus offset is 3 mm. The maximum possible focus offset is 4 mm.

Select **Microplate (flip image)** when scanning single microplates. When selected, images are flipped automatically after each scan so the origin (well A1) of the plate is in the upper left corner. (Microplate images must be flipped because the plate is scanned through the bottom.) Deselect **Microplate (flip image)** when scanning membranes, gels or mice.

The **Channels** check boxes is used to specify whether to detect fluorescence in the 700 channel, the 800 channel, or both. When both are selected, fluorescence from each dye is detected separately and stored in a separate image file.

The **Intensity** fields control the detector sensitivity and affect the band intensity on the image. If the intensity is set too high, the detector may saturate and produce white areas in the middle of intense bands/dots. (Saturated pixels are colored cyan if the image is being displayed as a grayscale image.) If the intensity is set too low, the image may not show any fluorescence even though there is adequate signal from the samples. LI-COR Presets use an intensity value of 5.0 for membranes, 8.0 for DNA gels and 5.0 for protein gels or microplates. These settings may need to be optimized for your gels or membranes due to the differing background fluorescence of various materials. Intensity values from 1 to 10 in increments of 0.5 can be chosen, as well as low intensity values L0.5 to L2.0. L2.0 is the lowest intensity value Odyssey can use for scanning.

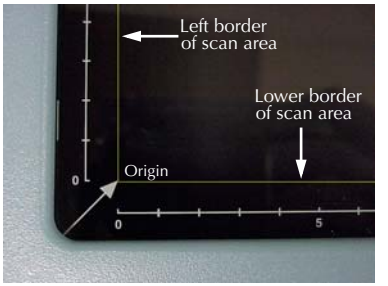
Scan Area parameters are used to specify the portion of the 25 x 25 centimeter scan surface to scan. The **Size** and **Origin** (cm) can be set by clicking and dragging a rectangle on the scan grid as shown below.



To reposition the scan area, click inside the red rectangle and drag the scan area to a new position. To resize the scan area, move the cursor over one of the red lines or corners until an arrow cursor is

displayed. With the arrow cursor displayed, click and drag to resize. To reset the scan area, click and drag a new rectangle on the scan grid, starting outside the current red rectangle. If necessary, double-click outside the current red rectangle to erase it before drawing a new one.

The tip of the arrow in the front left corner of the scanning surface on the Odyssey Imager corresponds to the **Origin** of X=0, Y=0 on the scan grid in the Scanner Console window.



See the Odyssey Operator's Manual for additional information on sample placement.

If the size and origin are known, the dimensions can be entered in the **Size** and **Origin** fields.

In general, it is best not to place the membrane or gel at the 0,0 position. The scan area drawn on the scan grid should always be larger than the membrane or gel so text annotations placed on the image during analysis will be displayed properly.

For low or medium resolution scans, make the scan area about 1 cm larger than the membrane or gel on all four sides. For example, if the membrane size is 5 x 5 cm, set the scan **Width** and **Height** to 7 cm, and set the Origin to 0,0. The membrane would then be placed at the 1 x 1 cm position on the scan surface.

Note: *After setting the scan area, check the file size at the bottom of the Scanner Console window to make sure the size is acceptable.*

Placing Samples on the Scan Surface

In general, it is easier to place the membrane or gel on the scan surface before drawing the scan area on the scan grid. If the sample is placed first, the 1 cm grid lines on the scan surface can be used to determine where to draw the scan area on the scan grid in the Scanner Console window.

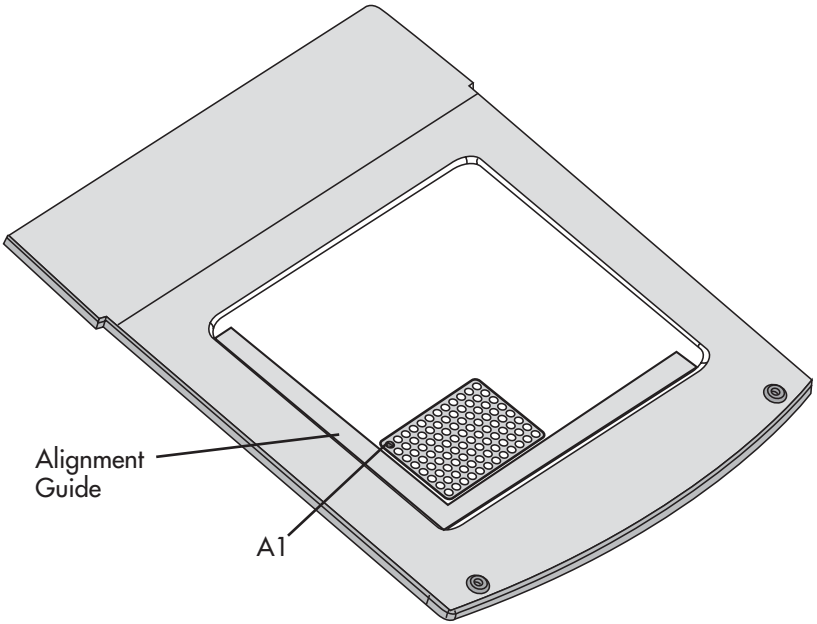
Membranes should be placed face down with the top of the membrane toward the front of the Odyssey Imager. (Orientation can be changed by flipping or rotating the image as needed.)

Tip: *Rectangular membranes (or gels) will scan faster if the long dimension of the membrane is oriented horizontally along the front border of the scan area. Placement in a vertical orientation requires the laser microscope to travel further and increases scan time.*

Consult the Operator's Manual and Odyssey Protocol pack inserts for tips on handling membranes and remember to touch the membrane only with a clean forceps.

Orienting a single microplate for a standard scan is somewhat different (scanning multiple microplates is described later in this chapter). A plastic microplate alignment guide is used to position the microplate at a known location on the scan surface. Push the guide into the lower left corner until it contacts the bezel surrounding the scan surface on both the front and left sides. Place the microplate on the scanning surface and slide it into position until it contacts both the front and left side of the alignment guide. The first well in the first row (A1) should be toward the back and left side of the alignment

guide as shown below. When the microplate is placed against the alignment guide, the scan size and origin parameters in the default microplate scan preset should work well.



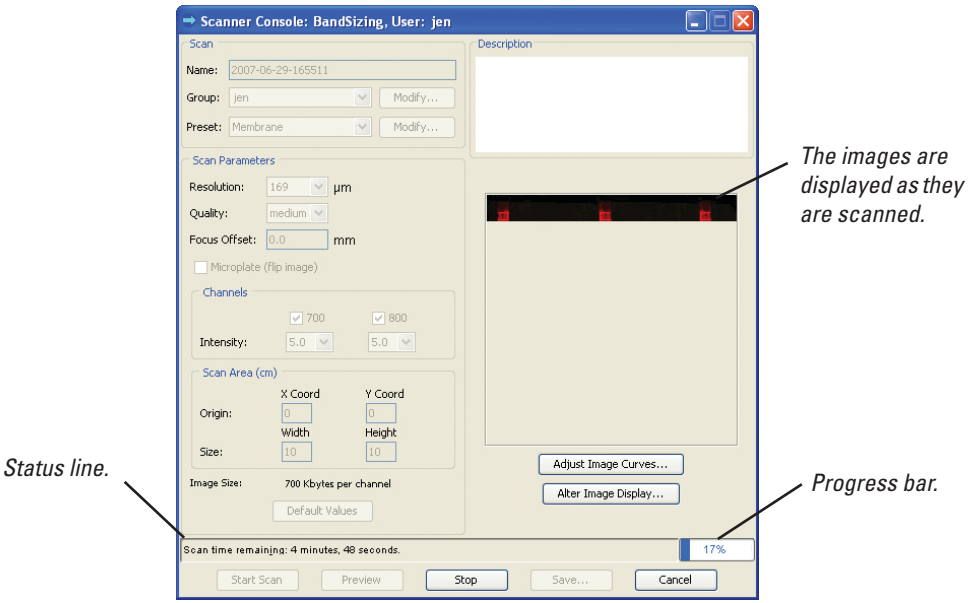
After placing the membrane, gel, or microplate on the scanning surface, close the lid on the Odyssey Imager.

For scanning mice with the MousePOD™ Accessory, consult the Odyssey *In vivo* Imaging Guide included with the MousePOD.

Starting a Standard Scan

To start a standard scan, click the **Start Scan** button in the Scanner Console to send the scan parameters to the Odyssey Imager and start the scan.

The images are displayed in real time in the area of the Scanner Console window where the scan grid was located.



At the bottom of the Scanner Console window, the status line indicates the time required to finish the scan. The progress bar indicates the percentage of the scan area that has been scanned. In the message area, the message "System Cooling" may be displayed initially, which indicates that the detectors in the laser microscope are being cooled to their operational temperature.

If no fluorescence is displayed where it is expected, click the **Alter Image Display** button to adjust the image (see Chapter 11). If bands are just dim, use brightness and contrast adjustments. If there are no bands, move the **Linear Manual Sensitivity** slider (auto adjustments off). The Adjust Image Curves window can also be used to make similar adjustments.

By default, the 700 and 800 channel images are shown overlaid. If the color scheme is the default red/green color scheme, areas that are yellow have intense fluorescence in both channels. To look at each channel separately during scanning, the Alter Image Display window can also be used to display one channel at a time.

If no fluorescence is visible, even after sensitivity adjustments, or if there is signal saturation (white pixels), cancel the scan (described below), and start the scan again using new values for the **Intensity** scan parameter in the Scanner Console. If fluorescence is too strong, use lower intensity values.

The scan ends automatically when the entire scan area has been scanned. As the images are collected, image files are created both on the hard disk of the Odyssey Imager and on the computer.

Stopping a Scan

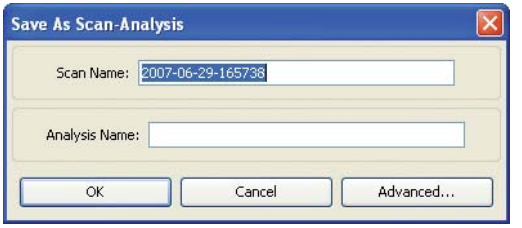
To finish a scan before automatic completion, click the **Stop** button in the Scanner Console window. The image files will be closed and saved, allowing the files to be analyzed.

To abandon a scan and not save the image files, click **Cancel** rather than **Stop**.

Completing the Scan

When the scan is complete, a reduced version of the image is shown on the scan grid and the **Save** button is activated so the scan on the Odyssey hard drive can be saved to the computer in an Odyssey project. To save the scan click **Save**. Alternatively, click **Close** to abandon the scan without saving any files.

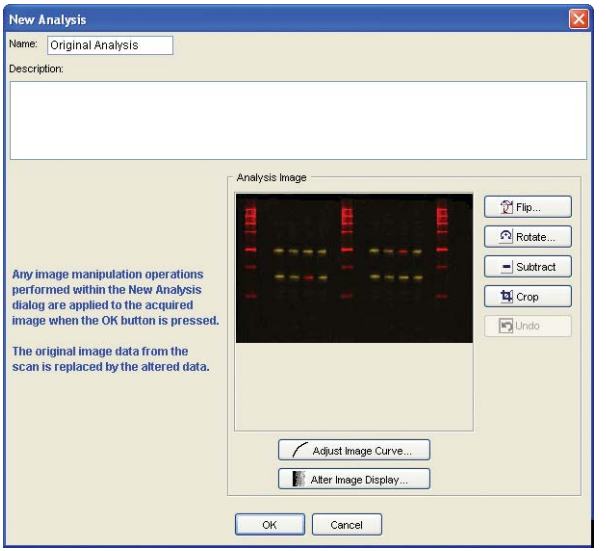
When **Save** is clicked, a dialog in displayed in which a new scan and analysis can be saved.



The scan name and analysis name are initially determined by the naming conventions specified in the Application settings (**Settings > Application** then **Naming Conventions**), but these names can be changed as needed. When **OK** is clicked, a scan folder is created in the current project and the TIFF image files are copied to the scan folder. An analysis with the specified name is also created in the scan folder. Both the scan and analysis are shown in the project directory in the main Odyssey window.

Occasionally, it may be necessary to manipulate the images before creating a new analysis. Click **Advanced** rather than **OK** in the Save As Scan dialog to manipulate the images before saving them.

When **Advanced** is clicked, the New Analysis window is opened.



Using the New Analysis window, a scan can be finished using one of two methods:

1) Enter an analysis name and click **OK** to save unaltered original images in a new analysis. To start analyzing the images, create another new analysis (Chapter 4) using copies of these original images.

2) Name the analysis and prepare the images for sizing or quantification using the buttons for flipping, cropping, rotating, background subtraction, or brightness and contrast adjustments. (See Chapter 3.) Then, click **OK** to create the new analysis and proceed with sizing or quantification.

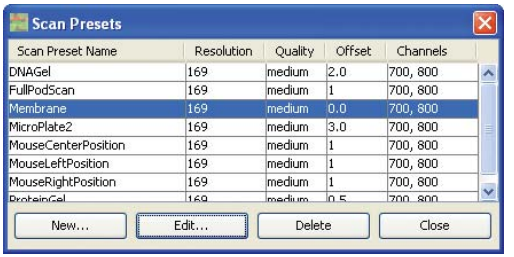
The first method uses more disk space, but is recommended for typical images since original copies of the scans are stored in the project.

If there is a problem with the scan and you do not want to analyze the images, click **Cancel** in the New Analysis window instead of **OK**. Clicking **Cancel** removes the new scan from the project that was open when the scan was started.

Creating and Editing Preset Parameters

After scanning a few of your own samples, you may want to create a set of Preset scan parameters or edit an existing Preset to match your scanning methods. The instructions below describe creating and editing Preset parameters for Odyssey Software. To create your own Presets in the Odyssey instrument, consult the Odyssey Operator's Manual.

The Preset parameters displayed in the Scanner Console window can be created or modified by choosing **Settings > Scan Presets**.



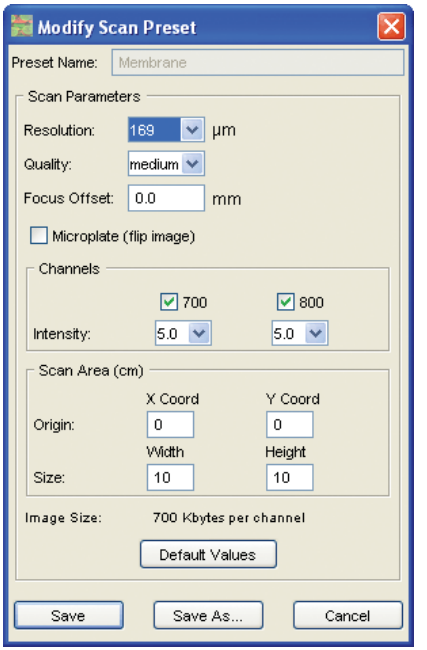
The Scan Presets window lists all presets. To edit a preset, select it from the list and click **Edit**. To create a new preset, either click **New** or edit an existing preset and click **Save As** in the Modify Scan Preset window.

Using the Modify Scan Preset Window

The Modify Scan Preset window is used to modify each of the scan parameters displayed in the Scanner Console Window. Scan parameters were defined earlier in this chapter.

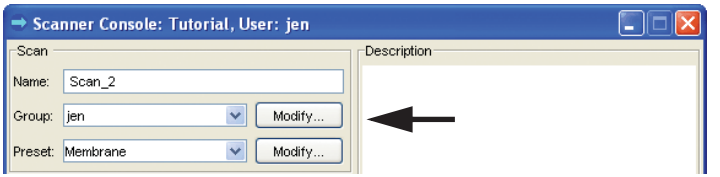
After you are done editing the scan parameters, check the image size to make sure it is under 14 MB and click **OK** if you just want to edit the Preset. If you want to create a new Preset with a different name, click **Save As** instead of **OK**.

If you make a mistake and want to set all values to stored factory default values, click the **Default Values** button. To abandon changes without saving them, click **Cancel**.



Note: *The status of the* ***Microplate (flip image)*** *parameter is ignored when a preset is used for a multi-plate scan. The images are always flipped for multi-plate scans.*

Tip: *The Modify Scan Preset window can also be opened by clicking* ***Modify*** *in the Scanner Console window as shown below. Note, however, that the* ***Scan Area*** *fields are not editable unless the Modify Scan Preset Window is opened from the* ***Settings*** *menu.*



Scanning Multiple Microplates

When scanning microplates, Odyssey can scan up to six microplates simultaneously. During a multi-plate scan, a separate scan and analysis is created for each microplate and they are added to the current project. For example, if there are six microplates, six scan files and their corresponding analysis files will be created. Odyssey relies on standard size plates being placed in a known location, so it is important to use the alignment guide and place the plates as described below. Use the following procedure to scan multiple microplates.

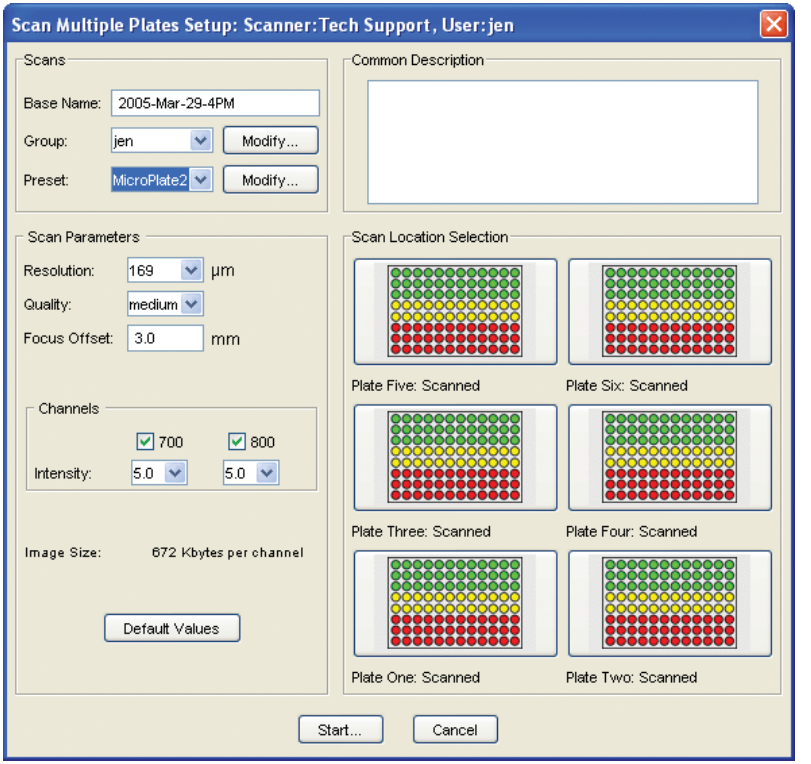
1) Create an empty new project or open an existing project in which to store the new scans (described earlier in this chapter).

2) Choose **File > Scan > Scan Multiple Plates**.

3) In the Scanner Login window, enter your **User Name** and **Password** (case sensitive) and click **OK**, if necessary.

4) In the Scan Multiple Plates Setup window, enter a base name for the scan or accept the default base name that is automatically entered according to the Multiple Scan settings (**Settings** menu).

Note: *Odyssey appends the base name with a sequential identifier for each microplate. In the window below, if 2005-Mar-29-4PM is the base name, Odyssey will automatically use 2005-Mar-29-4PM-1 as the scan name for the first plate, followed by 2005-Mar-29-4PM-2, etc. If another scan is started using the same base name, a letter is also appended. Thus, the first scan with a duplicate base name would be 2005-Mar-29-4PM-a-1, the second would be 2005-Mar-29-4PM-a-2, etc. The letter is incremented each time a new scan is initiated using a duplicate base name.*



5) Choose the scan **Group** in which to store the scan. (Use of scan groups in the Odyssey instrument is described earlier in this chapter.)

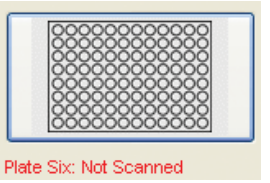
6) Enter a common **Description** (if any) that will be used for all microplates.

7) Choose a set of **Preset** scanner parameters designed for microplates (such as the default Microplate2 preset) and skip to the next step.

Alternatively, the individual scan parameters can be edited. The scan parameters are the same as those described earlier for standard

scans, with a few exceptions. First, the scan dimensions are not editable since a standard plate size is assumed. If non-standard plates are used, the scan size can be changed in the Multiple Scan settings. Second, set **Focus Offset** to 3.0 for standard microplates as recommended in the Odyssey Operator's Manual. The last difference is that the **Scan Multiple Plates** function automatically flips images to the correct orientation so the image has well A1 in the upper left corner. Presets for microplates can be created as described earlier in this chapter, but note that the scan size and origin are not used by the multiple scan software.

8) Set the number of plates to scan. If less than six plates are scanned, click the plate icons corresponding to empty locations in six-through-one order. Deselected plates appear as shown below.



Plates should be added to the Odyssey scanning surface in the order (1 through 6) shown in the multi-plate scan setup window. Consult the Operator's Manual for more details on placement of the scanning guide and plates.

9) Click **Start** to send the scan parameters to the Odyssey instrument and start the first scan.

The image from the first microplate is displayed in real time in the Scanner Console window. The name extension in the **Name** field indicates which plate is being scanned (-1, -2, etc.). The status line at the bottom of the Scanner Console window indicates the time required to finish scanning the current plate and a progress bar indicates how much of the plate area has been scanned.

If no fluorescence is displayed where it is expected, use the **Alter Image Display** or **Adjust Image Curves** button to adjust brightness and contrast. By default, the 700 and 800 channel images are shown overlaid. If the default red/green color scheme is being used, the areas that are yellow have intense fluorescence in both channels. If these adjustments do not display any fluorescence, you may need to start the scan again and scan with a different intensity value.

After the first scan is complete, an analysis containing the images is automatically created for the scan using the same name as the last analysis that was created. When the scan of the first microplate is complete, Odyssey automatically begins to scan the second plate and repeats the scan procedure until all designated plates have been scanned.

Note: *If the microplates are poorly centered in the images, try adjusting the grid template to match the grid to the wells in the image. If some wells are truncated or the grid is off the image, see Multiple Scan settings below for instructions on changing the scan offset to center the wells on the image for new scans.*

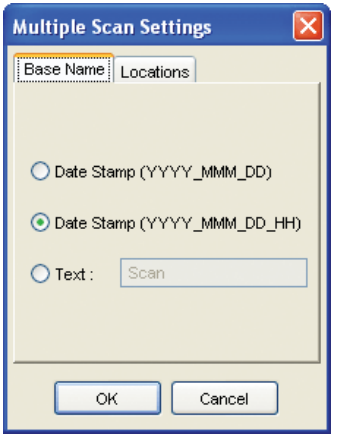
Stopping a Multi-plate Scan

To finish the scan of the current plate before automatic completion, click the **Stop** button. The image files will be closed and saved, and Odyssey will start the scan of the next plate (if any). To abandon a scan and not save the image files, click **Cancel** rather than **Stop**. Clicking **Cancel** also cancels all other plates from being scanned.

Multiple Scan Settings

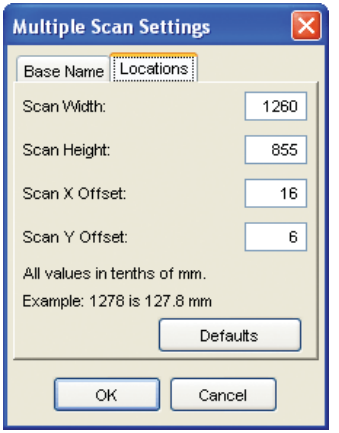
In the setup window for multi-plate scanning, a default base name is presented that is used to name all plate scans for a given series of scans. The default base name can be specified on **Base Name** tab in

the Multiple Scan Settings window. To open the Multiple Scan settings, choose **Settings > Multiple Scan**.



The default name can be set to the current date, the current date plus current hour, or user designated text.

In the setup window for multi-plate scanning, the user is not given the option of setting the scan size or XY offset. This is because the scan size is based on a standard microplate size. The scan size and XY offset for microplates recommended by LI-COR are shown below.



These settings are only a guide and every instrument may not produce images with the microplate exactly centered. If the microplate is not centered in the image, proceed with analysis and adjust the grid template to match the location of the wells. If the wells are truncated or if the grid is off the image, adjust the **Scan X Offset** and **Scan Y Offset** in the Multiple Scan settings to center the microplates in the images. Only integer numbers should be entered and units are 0.1 mm. For example, a **Scan X Offset** of "15" is an offset of 1.5 mm.

For non-standard plates scan sizes and offsets have to be determined experimentally. Start with the default values above and increase or decrease them as needed. Set the scan width and height inside the plate edges, but outside of the rectangle defining the well boundaries. Set the **Scan X Offset** to the distance along the X-axis from the right boundary of one scan to the left boundary of the next scan. Similarly, set the **Scan Y Offset** to the distance along the Y-axis from the upper boundary of one scan to the lower boundary of the next scan. If the scan size and offset are changed, you will also need to edit a grid template so the template matches the non-standard plate size.

Chapter 3: Creating a New Analysis

Overview

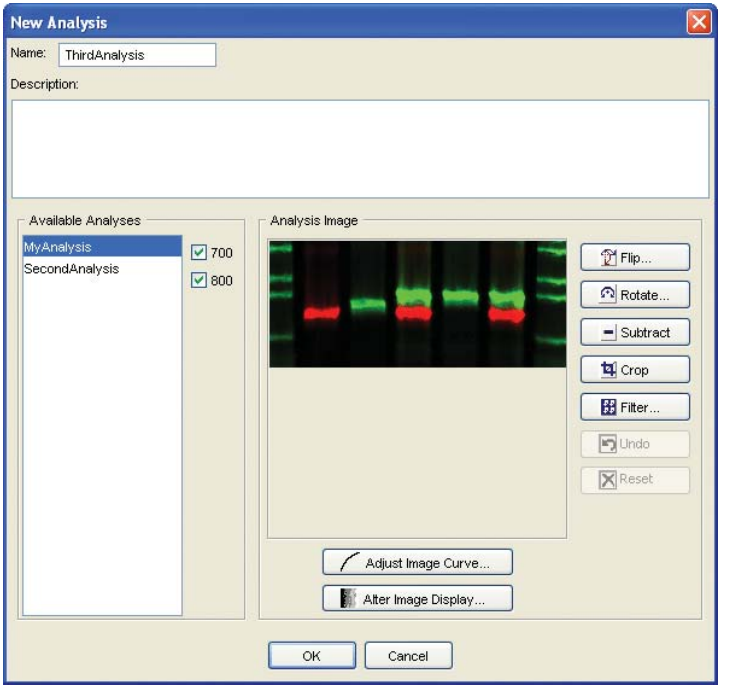
An analysis holds all the sizing or quantification data created when a scan is analyzed. For each scan, there can be many analyses. At the end of each new scan, a new analysis is named and saved along with the new scan, as described in Chapter 2.

For existing scans, a new analysis can be created by choosing **File > Analysis > New Analysis** to open the New Analysis window. Creating a new analysis makes copies of images from another analysis, allowing the images can be analyzed separately from the original analysis.

A new analysis may need to be created for a variety of reasons. For example, if the original image had multiple membranes scanned for different users, each user may want to independently analyze the portion of the image that is of interest to them. If the original image is in the wrong orientation, a new analysis can be created to flip or rotate the image.

Opening the New Analysis Window

To start a new analysis, open a project and click on the scan containing the images you want to copy into the new analysis. Next, choose **File > Analysis > New Analysis** to open the New Analysis window.



Naming the Analysis

The **Name** field is used to name the new analysis. In general it is best to use numbers, letters, underscore characters, or dashes. Do not use slashes, colons, or commas.

Entering a Description

Any text entered in the **Description** field can be included in reports after analysis is complete (Chapter 10). A list of the scan parameters and any alterations made to the image (cropping, etc.) is appended

to the end of the description. When the analysis is open, the description can be edited at any time by choosing **Edit > Analysis Description**.

Copying Images From Another Analysis

The **Available Analysis** list in the New Analysis window displays all images in the current scan that can be copied into the new analysis. When an analysis is selected from the list, a "thumbnail" view of the images that will be copied is shown in the **Analysis Image** section of the window.

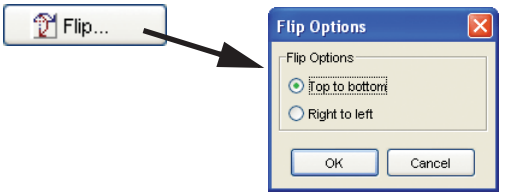
If there are two images in the selected analysis, both the **700** and **800** check boxes will be selected, indicating that both images will be imported. To copy only one of the images to the new analysis, deselect the image that is not wanted. See descriptions of the **Undo** and **Reset** buttons below for information about changing images after image manipulations have been performed.

Manipulating Images

The **Flip**, **Rotate**, **Subtract**, **Crop**, **Filter**, **Adjust Image Curve**, and **Alter Image Display** buttons in the New Analysis window are used to alter images before they are added to a new analysis. The **Undo** button cancels the last change made using the **Flip**, **Rotate**, **Subtract**, **Crop**, **Filter**, **Adjust Image Curves**, and **Alter Image Display** buttons. Odyssey has multiple undo capability, so clicking **Undo** multiple times steps backward through each change made since the New Analysis window was opened. To undo all changes in the New Analysis window, click **Reset**. After clicking **Reset** or using **Undo** to reverse all changes, the images selected in the Available Analysis panel can be changed.

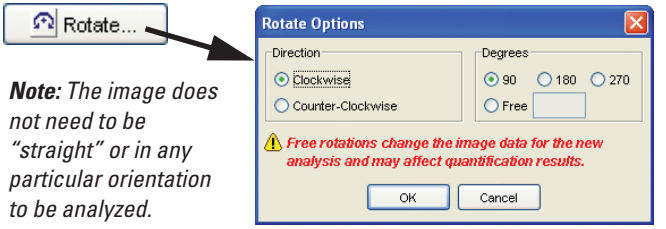
Flipping an Image

To flip an image vertically, click the **Flip** button and select **Top to Bottom** from the **Flip Options**. To flip an image horizontally, click the **Flip** button and select **Left to Right** from the **Flip Options**.



Rotating an Image

If the image is not in the desired orientation, click the **Rotate** button and use the Rotate Options window to rotate the image. If the image was scanned sideways or upside-down, use the **Direction** radio buttons to rotate the image either **Clockwise** or **Counter-Clockwise**, and then choose **90**, **180**, or **270** degrees of rotation.

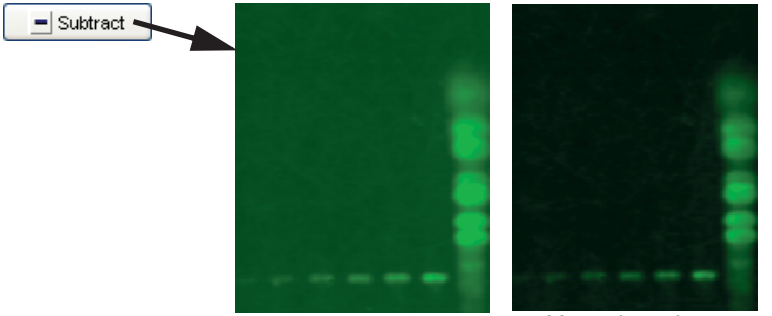


Note: *The image does not need to be "straight" or in any particular orientation to be analyzed.*

If the membrane was place on the scanner at a slight angle, it can be straightened using the free rotation option. First choose **Clockwise** or **Counter-Clockwise** rotation. Next, click **Free** and enter the desired rotation in degrees. Note, however, that *quantification results are changed* due to image interpolation when images are rotated to some angle that is not a multiple of 90 degrees.

Performing Background Subtraction

Background subtraction finds the minimum intensity value on the image and subtracts that intensity from all pixels in the image.



Before subtraction. After subtraction.

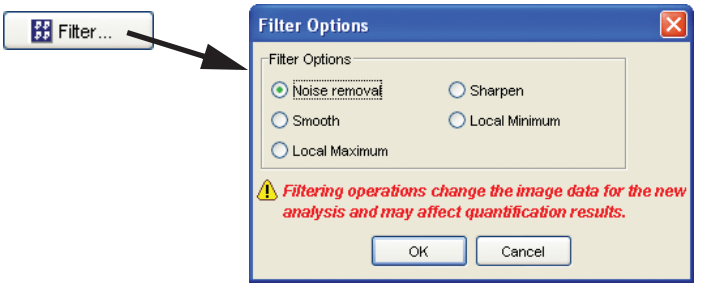
Cropping Images

To crop an image, draw a selection rectangle around the desired area of the image and click the **Crop** button. The selection rectangle is drawn by moving the cursor to one of the four corners of the crop area. Click and drag the cursor while holding down the left mouse button. Don't release the mouse button until the border of the selection area encloses the entire crop area.

IMPORTANT: Be careful not to crop too close to bands or other objects that will be analyzed. Leave some empty image around the edges. If the image is cropped too close to bands, it may be difficult to add text, lanes, or features near the edge of the window.

Using Image Filters

IMPORTANT: When quantifying bands/dots, none of the filters should be used since they change the image data, thereby invalidating the quantification results.



Each filter operation uses a 3 x 3 pixel convolution filter. A brief operational description of each filter is given below.

Noise Removal: The **Noise Removal** filter typically removes background speckles from an image. Noise removal calculates a median pixel value within the 3x3 filter region and replaces the current pixel value with the median. Noise removal is most noticeable where the median replaces pixel values that are much brighter or darker. For band sizing on "noisy" images, noise removal can improve band-finding accuracy because the lane profile will be smoother and have fewer noise peaks that might be identified as bands.

Smooth: The **Smooth** filter changes the current pixel value to the average of the pixels within the 3x3 filter region.

Local Maximum and Minimum: The **Local Maximum** filter replaces the current pixel value with the maximum pixel value within the 3x3 filter region. Similarly **Local Minimum** replaces the current pixel value with the minimum pixel value within the filter region. The **Local Maximum** filter reduces noise on images where there are background pixels that are much darker than surrounding pixels. The **Local Minimum** filter reduces noise on pixels where the background is fairly uniform except for some pixels that are much brighter than others.

Sharpen: The **Sharpen** filter multiplies each pixel in the region by sharpening coefficients and then sums the pixel values to create a new pixel value that replaces the current pixel value. The sharpening coefficients have been chosen so the edges of objects are enhanced.

Changing Brightness and Contrast

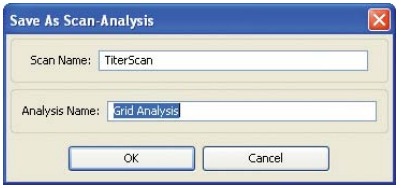
When the New Analysis window is opened, the image thumbnail is shown using the last-used brightness and contrast settings. For repetitive scans the last-used settings may work well. For other scans, brightness and contrast may need adjustment in order to see any fluorescence in the image. If fluorescence cannot be seen, changing brightness or contrast may be necessary to perform other image manipulations like cropping.

To change the brightness and contrast or to view only one of the two images, click either **Adjust Image Curve** or **Alter Image Display**. The operation of these two functions is described in Chapter 11.

Saving an Analysis

Any of the following methods can be used to save analysis files.

- Choose **File > Save** (or **Ctrl+s**) is used to save a project. All analyses with unsaved changes will be saved. Similarly, **File > Save As** saves all files and any unsaved changes from the current project to a new project folder.
- Choose **File > Analysis > Save Analysis** to save only the current analysis. All other analyses with changes will remain unsaved.
- Choose **File > Analysis > Save As New Analysis** to save the current analysis as a new analysis. Rename the analysis using the **Scan Name** field if the new analysis will be saved in the current scan. To save the analysis in a new scan, change the **Scan Name** (an analysis cannot be saved to an existing scan other than the current scan).



Note: *The* ***Save Analysis*** *and* ***Save As New Analysis*** *menu choices can also be accessed by right-clicking an analysis in the project directory tree and choosing the desired menu choice from the contextual popup menu.*

Deleting an Analysis

Caution: Use this function very carefully. The analysis is deleted immediately when this function is performed and the deleted analysis cannot be recovered.

To delete an analysis, click the analysis in the Scans view in the Odyssey window and choose **File > Analysis > Delete Analysis**.

Having More Than One Analysis Open

More than one analysis can be open at once, allowing images to be compared. When multiple image views are open, the **Window** menu can be used to switch between windows. The window name includes both the scan name and analysis name. Clicking the analysis name in the Scans view will also bring that analysis window to the front.

Chapter 4: Importing and Exporting Scans and Images

There are four methods to get scan images into a project in Odyssey software:

- Start a Scan: Images are automatically transferred to the computer when the scan is completed.
- Download a Scan: Scans started using one-button scanning on the Odyssey Imager can be retrieved from the instrument using the Download Scans function.
- Import a Scan: Scans stored in any Odyssey project can be imported into the current project.
- Import Images: Images that no longer have a scan file (*.scn) can be imported into the current project and stored under a new scan name.

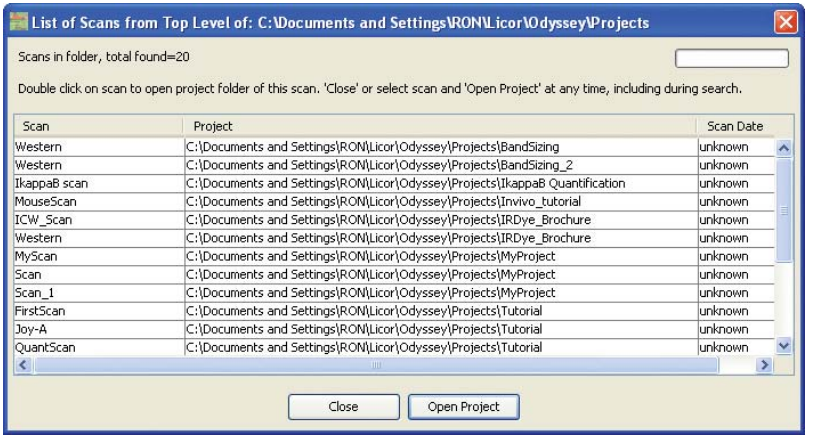
Starting a new scan is discussed in Chapter 2. The remaining three methods for importing images are discussed in this chapter. The following methods to export scans and save images to a new analysis are also discussed:

- Export Scans: Saves a copy of the current scan to any specified location.
- Flip or Rotate to New Analysis: Copies image files from the current analysis into a new analysis and flips or rotates the images in the new analysis.
- Crop to New Analysis: Crops image files in the current analysis and saves the cropped images in a new analysis.
- Crop to Multiple Images: Crops the current images using the scan area in a scan preset file and saves the cropped images to a new analysis.

Searching for Scans

Before importing or exporting a scan, you may need to search for it. Choose **File > Scan > Search for Scans** to generate a list of all scans in a folder or disk drive. A file dialog is displayed to choose the starting location for the search. If all Odyssey projects are kept in one folder, that folder may be a good starting place for the search. Entire drives can be searched by starting at C:, D:, etc. Note, however, starting at a specific folder takes less time than searching an entire drive.

During the search, any folder in the specified path is assumed to be a scan folder if *.scn and *.tif files are found in the folder. Scan folders found during the search are listed in a search results window.



Any scan in the search list can be opened by double clicking the appropriate row, or by clicking a row and clicking **Open Project**. If a search is still in progress, opening a scan stops the search, and opens the project containing the scan.

Initially, the table is sorted in ascending order by **Scan** name. Click the **Scan** header to switch to descending order. To sort by a different column, click the column header and click again to switch from ascending to descending order as needed.

To close the search window click **Close**. **Close** can also be clicked during a search to stop the search and close the window.

Downloading Scans

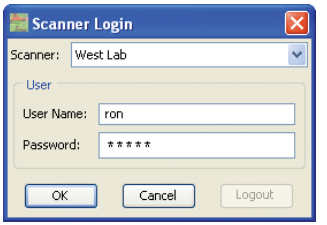
The Download Scan function is most commonly used to retrieve scans that were started from the Odyssey front panel keypad and stored in Odyssey's *public* scan group (see Odyssey Operator's Manual). By comparison, a scan started from within the Odyssey application software is downloaded to the computer automatically when the **Save** button is clicked at the end of the scan.

The Download Scan function can also be used to download another copy of a completed scan that is still stored on Odyssey.

Note: *Odyssey should not be used as a long-term storage device for scans. It is best to download scans immediately after they are finished.*

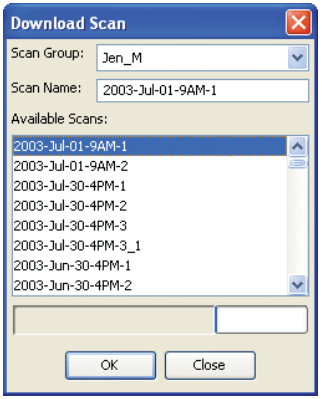
A project must be open before scans can be downloaded. Either a new or existing project can be used.

Start the download by choosing **File > Scan > Download Scan**.



In the Scanner Login window, select the scanner on which the files are stored and enter your user name and password.

After logging in, the Download Scan window is used to select a scan.



First, choose a scan group. The **Scan Group** drop-down list shows only scan groups that the current user belongs to. After choosing a scan group, select a scan from the **Available Scans** list. Chapter 12 describes the User Administration settings and how they are used to change access to scan groups, if needed.

Select a scan and click **OK**. To use a different scan name when the files are downloaded (to avoid duplicate scan names, etc.), enter a new name in the Scan Name field.

The selected scan will be downloaded into the current project. The scan can be analyzed by creating a new analysis (Chapter 3) after the files have finished downloading. A progress bar in the Download Scan window shows how much of the download has been completed.

Importing Scans

Occasionally, it may be necessary to copy a scan from one project to another to reorganize the projects. Choosing **File > Scan > Import Scan** imports scans from other projects into the current project. All analyses associated with the imported scan are also copied along with the scan.

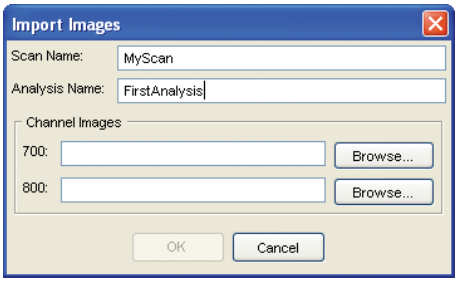
A project must be open before scans can be imported. After choosing **File > Scan > Import Scan**, use the standard file selection window to find the scan and open it. Only scan files (*.scn) are listed in the file selection window. Change the **Files of Type** field to **All Files** if you want to see all files. If a scan is not listed in a directory where it is expected to be, the scan file may have been deleted. In this case, choose **File > Scan > Import Images** instead of **Import Scan** (or download the scan from the Odyssey instrument).

Importing Images

During each scan, a scan file (*.scn) and one or two TIFF images (*.TIF) are created. If the original scan file has been lost or deleted, choosing **File > Scan > Import Images** can be used to import just the TIFF image files from a scan. **Import Images** can only import images from a local or network storage device, not from the Odyssey Imager.

Note: *When only images are imported, analysis files are not imported. The image resolution and scan remarks are also not imported since these data are in the scan file. Always use* ***Import Scan*** *if you want to import analysis files with the images.*

A project must be open before importing images. Images can be imported into an existing project or a new project can be created. To import images, choose **File > Scan > Import Images**.



The **Scan Name** and **Analysis Name** fields are used to name the new scan and analysis for the imported images. Images can only be imported into a new scan.

The **Browse** buttons next to the **700**- and **800**-channel image fields are used to browse for the image to import. After selecting one of the images, the image from the other channel is automatically found as long as the two image files are in the same directory. If the two image files are in different directories, click the button next to the other channel to browse for the second image.

After clicking **OK** to import the images, the new scan and analysis are displayed in the directory tree of the Scans view (Chapter 1) in the main Odyssey window.

Importing Images From Other Imaging Systems

The **Import Images** function is designed primarily for LI-COR TIFF images created by the Odyssey Imager. However, some 16-bit grayscale TIFF images from other imaging systems can also be imported. 8-bit grayscale TIFF images from common image editing programs cannot be imported.

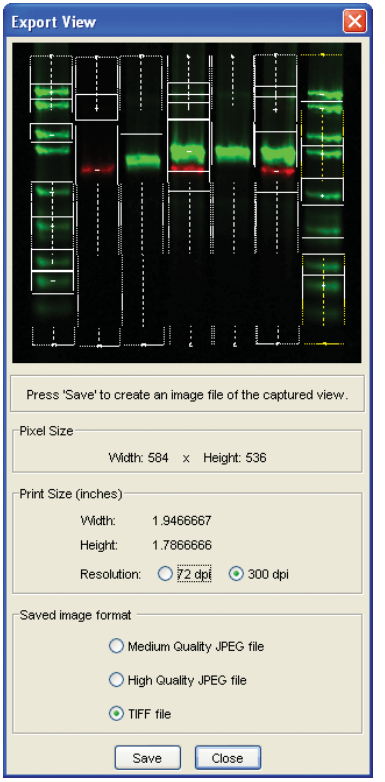
Exporting Images

Exporting an Image View

Choose **File > Export Image > Export Image View** to save the image to a JPEG or TIFF file. The portion of the image saved is the portion visible in the Image View. In the case of a pseudo color image, the pseudo color legend is appended to the right side of the image when it is exported. Before the image is saved, the image is shown in the Export View window and the file size is listed.

The image can be saved at a resolution of 72 dpi or 300 dpi. 72 dpi is generally preferred for electronic slide presentations and web applications. Some journals, however, require that images have a 300 dpi resolution or greater.

With either resolution, there is the same number of pixels in the file. The pixels stored in the file are exactly the same as the pixels shown on the computer monitor. To get the highest quality image for print applications, make the image as large as possible on screen before exporting it to a file. This will build an image file with as many image pixels as possible. (To get even more pixels in the file, use the Windows Control Panels to set the display resolution to a high level.)



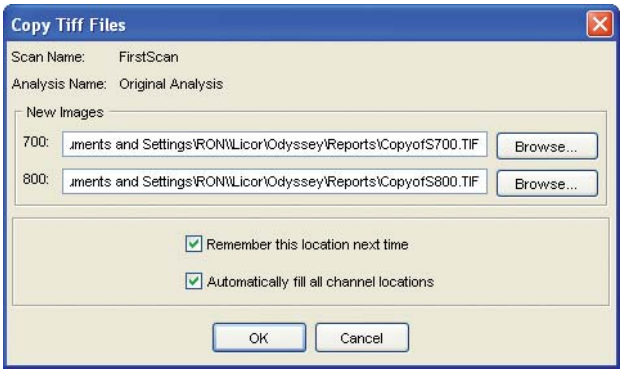
The image can be saved in three file formats. **TIFF** files will have the largest file size and the highest quality because there is no compression. The **High Quality JPEG File** setting uses a slight amount of JPEG compression to reduce file size, but still maintains very high image quality. **Medium Quality JPEG File** moderately compresses the image, which produces some noticeable compression artifacts in the image compared to an image that is not compressed. **Medium Quality JPEG** images are useful for e-mails due to the small file size and may be suitable for web or slide presentations if the image artifacts are acceptable. **Medium Quality JPEG** files are not recommended for print publication.

When the file is saved, all annotations currently displayed on the image will be saved in the JPEG file. Use the Application settings to select which annotations are displayed. Annotations can also be turned off or on by clicking  on the toolbar or by choosing **View > Hide Annotations**.

Note: *If the Application Settings (Image View Display) have been used to enlarge the text display area in order to prevent annotations from being truncated at the edge of the image, this extra text display area is included when an image view is exported (assuming the extra text area is visible in the Image View window).*

Exporting the TIFF Images

Choose **File > Export Image > Copy TIFF Files** to save copies of the 16-bit grayscale TIFF images to any location. 16-bit TIFF images can be used in Odyssey software and other analysis software that accepts 16-bit grayscale images. Commercial image editing programs (Adobe® Photoshop®, etc.) generally require 8-bit TIFF images (see *Exporting 8-bit Grayscale Images* below).



The **Browse** button opens a standard file selection window. If **Automatically fill all Channel Locations** is selected, the path for the second channel is automatically updated when the path for the first

channel is chosen using the **Browse** button. To always save the copied images in the same location, select **Remember this Location Next Time**.

Exporting 8-bit Grayscale Images

Choose **File > Export Image > 8-bit Grayscale TIFF** to export 8-bit grayscale TIFF images rather than the 16-bit grayscale images scanned by the Odyssey System. 8-bit grayscale images can be used with commercial image editing software and presentation software such as Microsoft® PowerPoint®. The Create Grayscale Tiff window works as described above for copying TIFF files.

Note: *Quantitative analysis should not be performed on exported 8-bit images.*

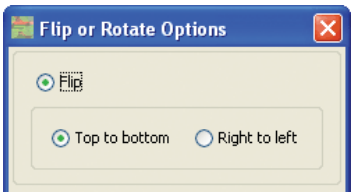
Exporting Colorized TIFF Files

Choose **File > Export Image > Colorized TIFF File** to save a single TIFF image file that contains both the 700- and 800-channel images. The images are colorized and overlaid, resulting in the same appearance as the images in the Odyssey window. **Colorized TIFF File** export can also be used for single images in pseudo color format. Annotations and other markers are not included -- only the image data are saved.

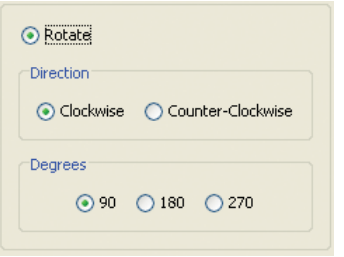
Flip or Rotate to New Analysis

Choose **File > Analysis > Flip or Rotate to New Analysis** to copy image files from the current analysis into a new analysis and flip or rotate the images in the new analysis.

To flip the images, select **Flip** and select the direction to flip the images (top-to-bottom or right-to-left).

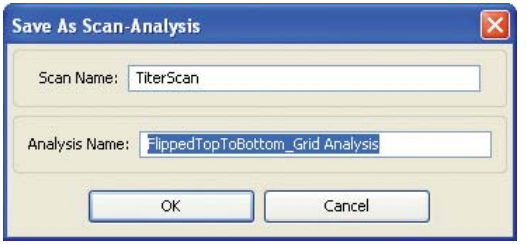


To rotate the images, select **Rotate**, and select the direction and degrees to rotate the images.



Note: *To rotate the images by an increment other than 90 degrees, create a new analysis and rotate the images in the New Analysis window (Chapter 3).*

After clicking **OK**, the Save As window opens with the current scan name and a proposed analysis name already filled in. To save the new analysis to the current scan, click **OK** or edit the analysis name and then click **OK**.

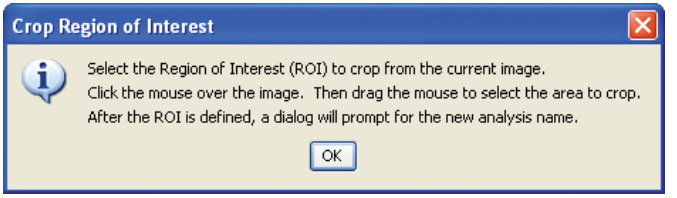


To save the new analysis in a new scan, enter a new **Scan Name** and **Analysis Name** (optional) before clicking **OK**. An analysis cannot be saved to an existing scan other than the current scan.

Crop to New Analysis

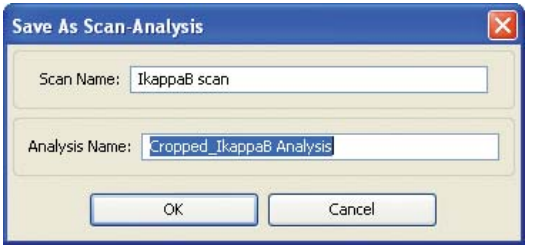
Choose **File > Analysis > Crop to New Analysis** to crop image files in the current analysis and save the cropped images in a new analysis.

A reminder is displayed that explains how to crop a region of interest.



After clicking **OK**, move the mouse cursor to one corner of the region of interest on the image. Click and hold down the left mouse button while dragging the cursor to the opposite corner if the region of interest. Release the mouse button.

To save cropped images in the current scan, click **OK** or edit the analysis name and then click **OK**. To save the images in a new scan and analysis, change the **Scan Name** and **Analysis Name** (optional) before clicking **OK**. An analysis cannot be saved to any scan other than the current scan or a new scan.



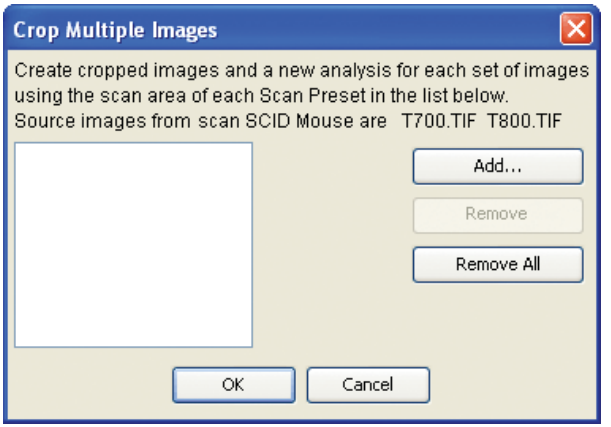
Crop to Multiple Images

Choose **File > Scan > Crop to Multiple Images** to crop the current images using the scan area in a scan preset file (Chapter 2) and save the cropped images to a new analysis. If necessary, create and save scan preset(s) with the scan area equal to the cropped size of the images.

Example: Suppose you are using the Odyssey MousePod, which has three mouse positions. The original scan was collected with the FullPodScan preset, which creates one image that includes all three mouse positions. Now, however, it would be more convenient to analyze each mouse separately. The **Crop to Multiple Images** function can be used to create three new analyses using the scan presets for the left, middle and right mouse positions.

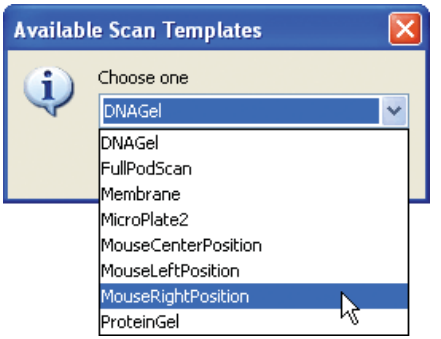
Here is how the three new analyses can be created.

1) Begin by choosing **File > Scan > Crop to Multiple Images**.

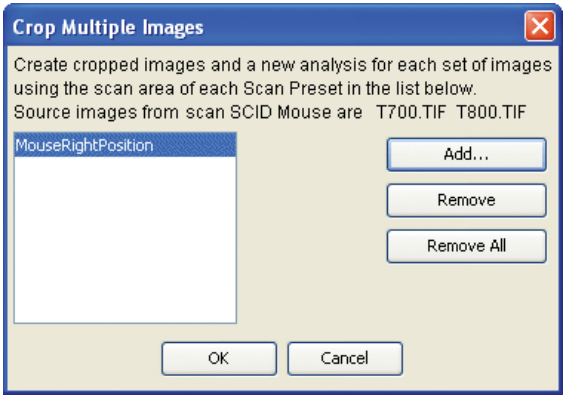


2) Click **Add** to add a scan preset to the list of scan presets.

3) Choose a scan preset that has a scan area in the correct position to use as a cropping template (the MouseRightPosition preset in our example).



4) Click **OK**.



5) Repeat steps 2 - 4 for the center (MouseCenterPosition) and left (MouseLeftPosition) mouse positions. If there are any other scan presets you would like to use, add them to the list. The current images can be cropped many different ways by choosing additional scan presets.

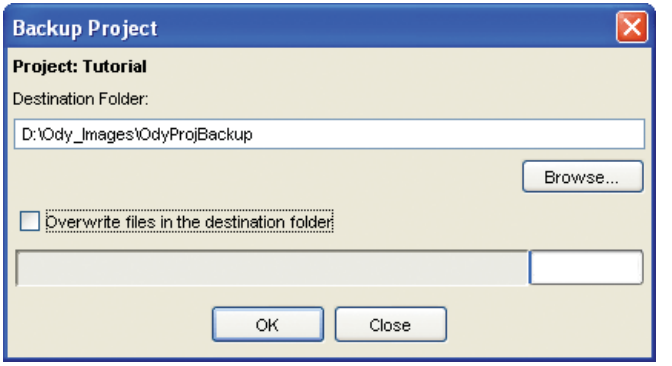
6) Click **OK** to crop the current images and save each of the three cropped images in new analysis files.

Exporting Scans

Choose **File > Scan > Export Scan** to save a copy of the current scan to any location (network, local, etc.) specified in the Select Destination Folder window. The scan folder and all of its contents will be copied, including all analysis files.

Project Backup

For archival or safety purposes, the project that is currently open can be backed up to any destination by choosing **File > Backup Project**. Click **Browse** to open a standard file selection window or enter a path for the destination folder. Files can be stored on any local or network drive.



If you intend to frequently back up projects to the same directory after making changes, click **Overwrite Files in the Destination Folder** so the older copy is overwritten by the new version.

Chapter 5: Creating Lanes and Finding Bands

Lanes can be created by several methods. Typically, lanes are created one at a time by selecting the lane tool, clicking at the top-center of a lane, and then double-clicking at the bottom-center. Lane boundaries are drawn according to the lane width specified in the Application settings. Immediately after a lane is created, all bands are identified and marked with a band marker.

Before You Begin...

Creating or Opening an Analysis

An analysis must be open in order to add lanes to scanned images. For new scans, a new analysis is created as described in Chapter 2. For an existing analysis, double click the analysis in the navigation tree to open it.

Single Channel Display vs. Overlaid Image Channels

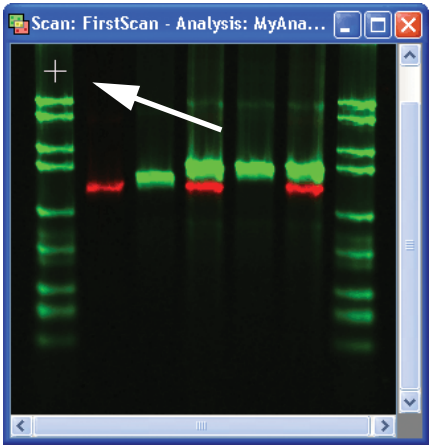
Scans with two images can be displayed in single channel mode, or with both image channels overlaid as a composite image. Lanes are usually added with images overlaid because lanes are added to both images simultaneously. Lanes can also be added to each image channel separately, if desired. After adding lanes, each image must be analyzed separately in single channel mode (Chapters 6 and 8).

Before creating lanes, click  on the tool bar until both images are overlaid in a composite image (assuming the scan has two images).

Creating the First Lane

To create a lane, choose **Lane > Add Lane** or click the (add lane) tool on the toolbar.

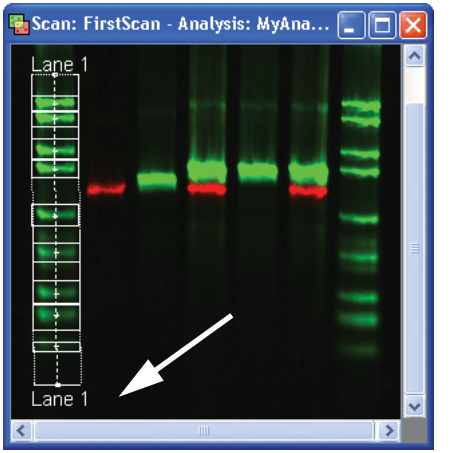
Click at the top of the lane you want to add. The mouse pointer should be horizontally centered in the lane.



Finding Straight Lanes

For straight lanes (vertical or slanted), center the mouse under the bottom of the lane and *double-click*. The lane will be added using the lane width specified in the Application Setting. Bands in the lane will be found and enclosed by band markers.

Lanes can be added from top-to-bottom or bottom-to-top.



TIP: *If you click and hold down the left mouse button at the top of the lane and drag the mouse downward to the bottom of the lane, a white dashed line extends from the point of the mouse click to the current cursor position. The white line indicates where the centerline of the lane will be. The white line in useful for creating straight lanes. When the white line is in the correct position, release the mouse button and double-click where the endpoint of the line should be.*

Finding Curved Lanes

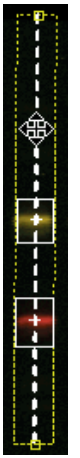
For curved lanes, begin by clicking at the top-center of the lane as usual. Next, add inflection points by single-clicking in the center of the lane at various points in the lane. Single-clicking creates a multi-segment lane line consisting of all the points that were clicked. Finish the curved lane by double-clicking the last point. The more a lane curves, the more inflection points will be required to make the lane conform to the image.

After the first lane is added, additional lanes can be added using the same technique. For greater efficiency, however, lanes can be copied and pasted, or multiple lanes can be added at once. Both techniques are described later in this chapter.

Moving and Resizing Lanes

Moving Lanes

To move a lane with image channels overlaid, click the lane to select it (changes color). Move the cursor into the middle of the lane until the cursor shows arrows in all directions. With the "all arrows" cursor displayed, click and drag the lane to a new position.

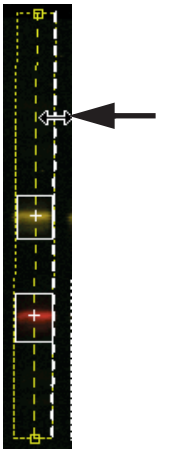


If lanes are moved in single channel mode rather than with channels overlaid, an error message will be displayed if the lanes are still linked.

Linked Lanes

When lanes are added to overlaid images, the lanes on both channels are "linked". In other words, when the images are overlaid and the lane position or width is changed, lanes on both the 700 and 800 channel images will be changed. Certain operations, such as trying to change lane width in single channel mode, will "unlink" the lanes on the two images. Band markers are never linked and can never be edited with channels overlaid. Bands in the lanes are found independently and automatically on both images immediately after lanes are marked.

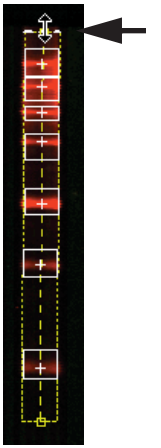
Changing Lane Width



To change lane width, click the lane to select it. Move the cursor over the left or right boundary of the lane until the cursor turns to a right-left arrow cursor. With the right-left cursor displayed, click and drag one side of the lane boundary (the centerline and opposite boundary are not moved).

To symmetrically change both left and right lane boundaries, click the Properties button ( ), and select the **Symmetric Left and Right Boundaries** check box.

Changing Lane Height



To change lane height, click the lane to select it. Move the cursor over the top or bottom boundary line of the lane until the cursor turns to an up-down arrow cursor. With the up-down cursor displayed, click and drag the lane boundary to a new position.

Changing Lane Shape

If one of the end points is not centered in the lane and needs to be moved to match the shape of the lane on the image, start by clicking the end point to select it. When the end point changes color and is

surrounded by the white selection box as shown below, the end point can be clicked and dragged to a new location.

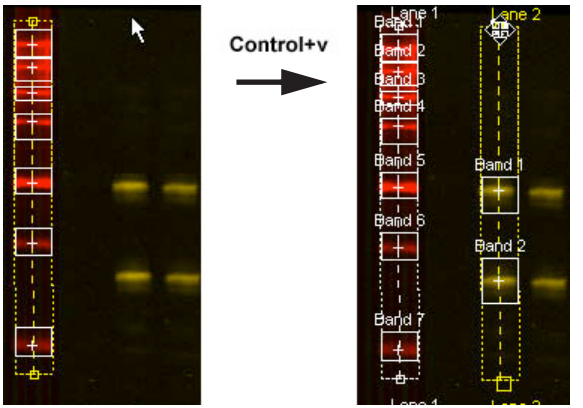
Straight lanes can be made vertical or slanted by moving one of the two end-points. Curved lanes can be reshaped using the same click-and-drag method on any point in the lane.

Copying and Pasting Lanes

After creating the first lane, additional lanes can be added by copying and pasting lanes. This method, however, is useful only if the other lanes on the image have a similar shape.

Copy a lane by selecting it and pressing **Ctrl+c** on the keyboard.

Lanes are pasted at the cursor position, unless the **Paste Special** command is used. Move the mouse cursor to a point on the image that is centered and at the top of an unmarked lane (shown below).



Press **Ctrl+v** on the keyboard to paste a new lane at the position of the mouse pointer. Bands in the new lane are found automatically.

Note: *Cut, Copy, Paste, and Undo are also available as tools on the left-side toolbar.*

Copying Multiple Lanes

Multiple lanes can be selected by holding down the **Control** key and clicking each lane, or by clicking and dragging a selection box around the desired lanes. **Ctrl+c** and **Ctrl+v** can then be used to copy and paste the lanes.

Using the Paste Special Command

The **Paste Special** choice on the **Edit** menu can also be used to paste lanes that have been copied, but the result is slightly different than the **Paste** choice (**Ctrl+v**). **Paste Special** pastes lanes in the exact same coordinate position as the lanes that were copied. **Paste Special** is generally used to copy lanes between images. Lanes can be copied between images in the same analysis or between images in different analyses.

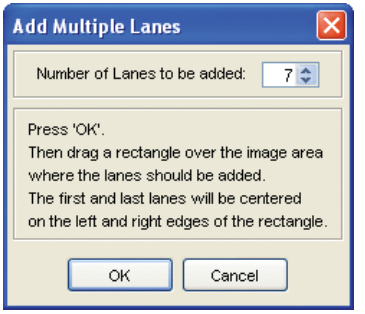
Deleting Lanes

Lanes can be selected and then deleted by clicking the (✕) button. If image channels are overlaid when a lane is deleted, lanes in both image channels are deleted.

Creating Multiple Lanes

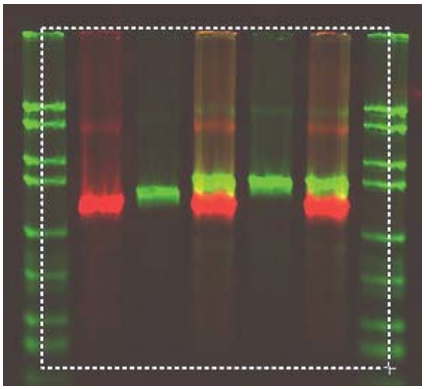
The Add Multiple Lanes tool can be used any time there are evenly spaced lanes on the image. This tool does not "find" lanes, but instead creates the specified number of lanes and spreads them evenly over the area marked by the user. Only vertical, straight lanes can be created. Before starting, check the default lane width using the Application settings as described at the end of this chapter.

Start by clicking the multi-lane tool ( ) or choosing **Lane > Add Multiple Lanes**. In the Add Multiple Lanes window, enter the number of lanes to create and click **OK**.



After clicking **OK**, click and drag a rectangle over the image area where the lanes should be added. Start in the center (horizontally) of the first lane, at a height that will correspond to the upper lane

boundaries. Release the mouse button when the right side of the rectangle is centered in the right-most lane and at the vertical position where the lower boundary of the lanes should be.



When the mouse button is released, the specified number of lanes is placed on the image. Lane width is determined by the Application settings. Bands are automatically found and enclosed by a band marker after the lanes are created.

Verifying Band Finding

After lanes are created and bands are found automatically, each lane should be visually checked to make sure all bands are marked with band markers and that the markers are in the correct position.

First, switch to single channel mode by clicking  . Visually check and make sure there is a band marker (rectangle) surrounding every band, and that there are no extra bands identified in the lane.

Too Many Bands

If there are too many bands in a lane, bands can be deleted by clicking an unwanted band marker to select it and clicking ✕ on the toolbar. Legitimate bands should have the '+' symbol centered over a band. False bands will be centered over empty image.

Not Enough Bands

Bands are added by clicking ( ) on the toolbar, centering the mouse pointer over the band on the image and clicking the left mouse button. Repeat the procedure for additional missing bands.

Fine-Tuning Band Finding

If there are consistently too many or too few bands on all images, the default band finding threshold can be adjusted in the Application settings as described later in this chapter.

Verifying Band Markers Are Centered

For both molecular weight sizing and quantification, it is important that band markers are centered in the bands. Visually check each band to make sure the '+' symbol in the band marker is centered in the band. All bands in a lane can be checked at once using the Lane Profile window as described later in this chapter.

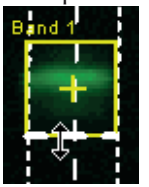
If a band marker is not centered it should be moved by selecting the band marker and moving the cursor to the center of the band marker. When the cursor changes to an all-arrows cursor, click-and-drag vertically to move the band marker until the '+' symbol is centered.

Verifying Bands Are Fully Enclosed

If the bands are going to be quantified, make sure each band is fully enclosed by the band marker and that each band marker is properly placed for the background calculation method in use (see Chapter 8). In general, the band marker lines should be over empty background image, if possible, and not touch any pixels that represent band fluorescence.

For MW sizing, placement of the band marker boundaries is not as important as long as only one band is enclosed in each band marker. If bands are not fully enclosed, sizing will be unaffected.

To change the size of a band marker, select the band marker and move the cursor over the top or bottom boundary line. When the cursor changes to an up-down arrow cursor, click-and-drag to move the boundary to the desired position.

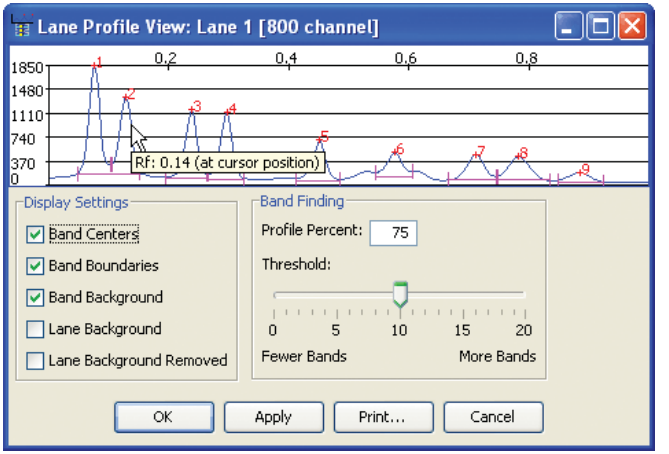


Re-finding Bands

If mistakes are made in band editing, the **Undo** choice on the **Edit** menu can sequentially undo each edit. At times, however, it may be easier to just start over. To re-find bands using Odyssey's automatic software, select a lane or lanes and then choose **Lane > Refind Bands**. Use this function carefully! All band data in the selected lanes are deleted before bands are found again.

Using the Lane Profile Window

A fluorescence profile for a selected lane can be displayed by clicking on the toolbar or choosing **Lane > Lane Profile**.



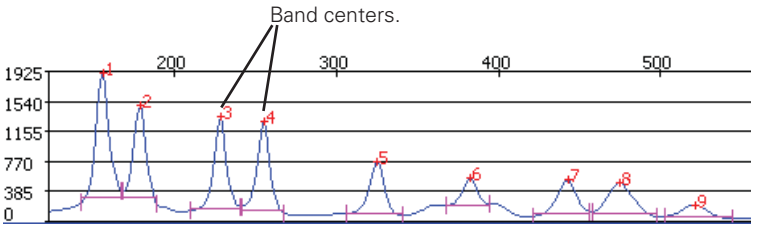
The title bar of the lane profile indicates which lane is profiled and the image on which the lane is located. If lane profiles are opened with channels overlaid, two profiles are opened – one for the lane in the 700 channel and one for the lane in the 800 channel. Similarly, if more than one lane is selected, profiles will be opened for all selected lanes. All lane profile windows can be closed by choosing **Window > Close Lane Profiles**.

The lane profile can be used for the following tasks:

- To check band finding
- To verify that band markers are centered on the fluorescence peaks
- To change band finding threshold and re-find bands
- To view a graphical display of background fluorescence in the lane
- To view the fluorescence peaks with background fluorescence removed

Understanding the Lane Profile

The X-axis of the lane profile represents the vertical position in the lane. The scale for the X-axis is shown above the graph and units are user selectable in the Application settings for lanes.



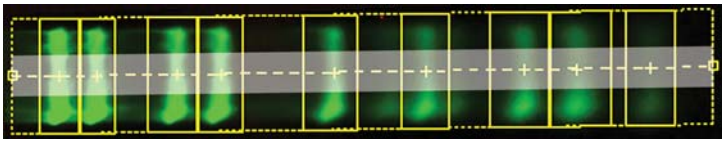
Y-Axis
Average intensity of all pixels within the lane profile width for a given row of pixels.

Lane Top

X-Axis
Molecular weight or band migration value. Migration in pixels is shown.

Lane Bottom

Lane profile width (percentage from settings).



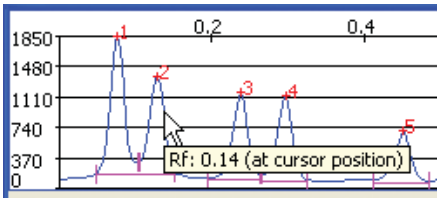
Lane rotated 90° counter-clockwise for illustration.

Units for the X-axis can be:

- **Pixel Location** (shown): distance in pixels from the image to the band center in pixels.

- **Relative Mobility**: Relative distance from the top of the lane. The top of the lane (left side of lane profile) is assigned a value of zero and the bottom of the lane (right side of lane profile) is assigned a value of one.
- **Size Standard**: Molecular weight calculated using sized standards currently applied to the image. Units are those previously assigned to the size standard when it was created.

The cursor position on the X-axis is shown in the cursor tool tip when the cursor is stopped over the graph. Below, the X-axis is set to display relative mobility and the cursor position is 0.14.

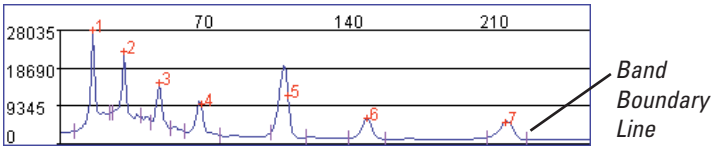


Each Y-axis value in the lane profile is the average intensity of all pixels within the profile width for a given row of pixels in the lane. The profile width of the lane is determined by the Application settings for lanes. Profile width is the percentage of the lane width (centered in the lane) used to calculate the lane profile. For a lane that is 40 pixels wide and a profile width setting of 75%, the pixel intensity values are averaged for the 15 pixels on both sides the lane's vertical centerline (30 pixels total). The average intensity is then plotted on the lane profile to represent that row of pixels. This is repeated for each row of pixels in the lane until the profile is complete.

Displaying Band Centers

With **Band Centers** selected in the Lane Profile window, the band number is displayed on each fluorescence peak along with a '+' that indicates where the band marker is centered. Displaying band centers is an easy way to make sure each band marker is centered over the

band on the image. In the lane profile below, band 5 is not properly centered since the center marker is not on the apex of the peak. If the image for band 5 were examined, the center of the band marker would not be centered in the fluorescence on the image and should be moved so the correct size will be assigned.

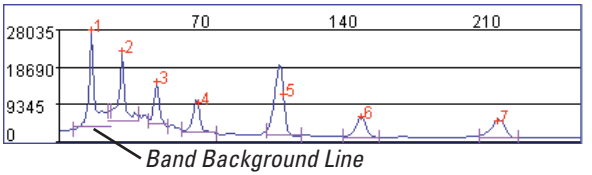


Displaying Band Boundaries

When **Band Boundaries** is selected in the Lane Profile window, vertical magenta lines are placed on the profile at the beginning and end of each band, as shown above. These lines correspond to the upper and lower sides of the band marker on the image. If lane finding identifies too many bands, band boundaries are useful for identifying false bands. False bands are usually centered on some small noise peak in the fluorescence curve rather than on a fluorescence peak. If a band marker is too small, it will be apparent in the lane profile because the band boundaries will be on the sides of a peak rather than down in the image "noise" between peaks.

Displaying Band Background Fluorescence

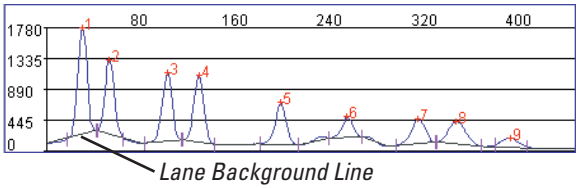
With **Band Background** selected, the calculated band background is displayed as a horizontal line between the band boundary lines.



Displaying the band background line can be useful in determining whether to use the band background or lane background as the background calculation method during quantification (see Chapter 8).

Displaying Lane Background Fluorescence

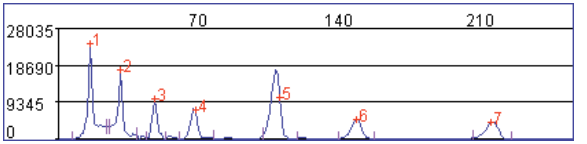
When **Lane Background** is selected, a profile for the lane background is displayed beneath the lane profile.



The lane background line is calculated by selecting appropriate minima points from the lane profile. Displaying the band background line can be useful in determining whether to use the band background or lane background as the background calculation method during quantification (see Chapter 8).

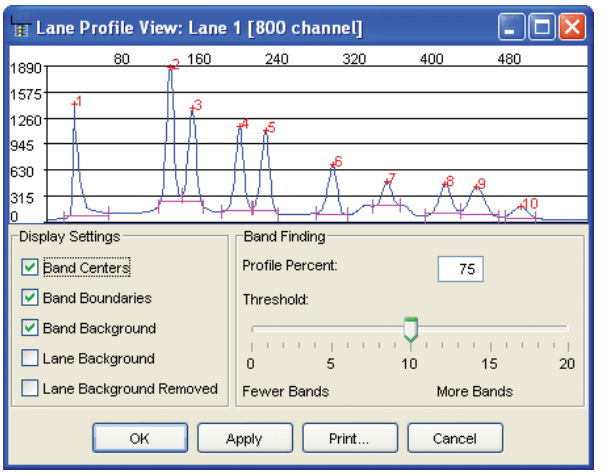
Displaying Lane Profiles With Background Fluorescence Removed

When **Lane Background Removed** is selected in the Lane Profile window, the lane profile is displayed with the lane background subtracted.



Controlling Band Finding Using the Lane Profile Window

The **Threshold** slider in the Lane Profile window is used to control the band finding software in order to more accurately find the correct number of bands in a lane.



As the **Threshold** slider is moved, band finding is changed in real time and bands are added or deleted on the lane profile as the slider moves. If the threshold is set too high, too many bands will be found when new lanes are pasted or created. If the threshold is set too low, too few bands will be found. The correct threshold setting for a given lane is the threshold setting at which the number of bands matches the number of fluorescence peaks. After adjusting the threshold, click **Apply** to apply the changes to the lane on the image. To optimize the default threshold setting, observe typical threshold settings on a variety of scans and then change the Application settings (described below) to match the typical threshold.

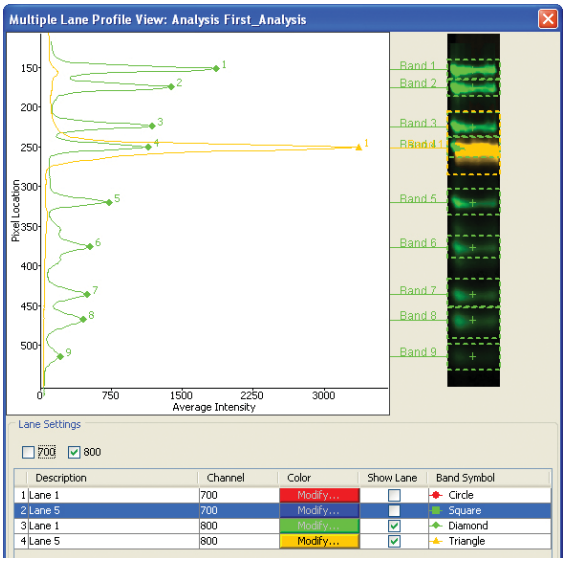
On images with a lot of background noise, the profile may be very jagged with noise peaks on the sides of larger fluorescence peaks. These noise peaks can cause inaccuracy in band finding even though

the band finding threshold is set correctly. If bands are not going to be quantified, the Noise Removal or Smoothing filters can be used to reduce noise peaks and improve band finding accuracy. These filters are available when starting a new analysis (Chapter 3).

Comparing Lane Profiles

To compare lane profiles for two or more lanes, select the lanes (**Ctrl + click**) and choose **Lane > Multi Lane Profile**. The window below shows a molecular weight standard lane (lane 1) and a sample lane (lane 5). The lane profiles show that band 1 in lane 5 is at the same pixel location as band 4 in the molecular weight standards lane.

A composite image of all the selected lanes is shown to the right of the profiles. Bands are colored with their respective lane colors. Band labels to the left of the image are also colored with the lane color. Lane colors can be changed by clicking the **Modify** button in the **Color** column in the lane table at the bottom of the window.



Clicking in the appropriate cell in the **Band Symbol** column of the lane table and selecting a symbol from the list of symbols changes the band symbol used in the lane profile.

To simplify the lane profiles, all lanes for a given channel can be turned off by deselecting either the **700** or **800** check box. Individual lanes can be turned off using the check boxes in the **Show Lane** column.

Normalizing Bands in Lanes

The sample normalization function allows one image channel to be used as a reference to normalize bands in lanes in the second image channel.

Normalization Procedure

1) With image channels overlaid, add lanes.

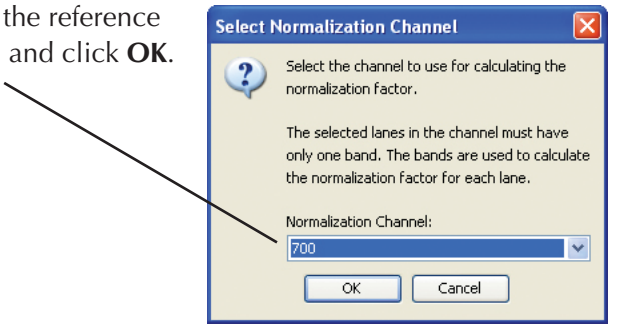
2) Click  and then  as needed until the channel is displayed that will be used as the reference channel.

3) Examine each lane in the reference channel image and make sure there is only one band per lane. Delete any surplus band markers.

4) Click  to display both images again and select (**Ctrl + click**) the lanes to normalize.

5) Choose **Lane > Sample Normalization**.

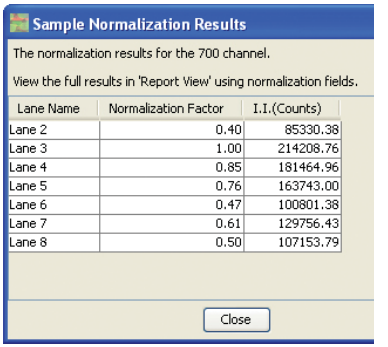
6) Choose the reference channel and click **OK**.



The reference channel is used to calculate a normalization factor for each selected lane. The band having the highest integrated intensity is assigned a normalization factor of 1.0. This reference band is then used to calculate the normalization factor for all other selected lanes by dividing the band integrated intensity by the integrated intensity of the reference band. On the sample image, a band's normalized integrated intensity is calculated by

Band Integrated Intensity / Lane Normalization Factor

After the calculations are complete, the normalization factors are shown in a table. The table shown to the right indicates that lane 3 contained the reference band and was assigned a normalization factor of 1.0. Both the normalization factor and the normalized integrated intensity can be included in reports or exported data.



Lane Name	Normalization Factor	I.I.(Counts)
Lane 2	0.40	85330.38
Lane 3	1.00	214208.76
Lane 4	0.85	181464.96
Lane 5	0.76	163743.00
Lane 6	0.47	100801.38
Lane 7	0.61	129756.43
Lane 8	0.50	107153.79

Creating and Using Lane Templates

Lane templates record lanes from one analysis so they can be applied to another analysis with similar lane patterns. Lane templates work well in repetitive experiments, such as those in which the LI-COR MPX Blotter is used to produce images with consistent lane and band patterns. Lane templates record all sample lanes, band markers, molecular weight standard lanes, and the applied set of molecular weights.

Saving a Lane Template

Start by doing all of the tasks below.

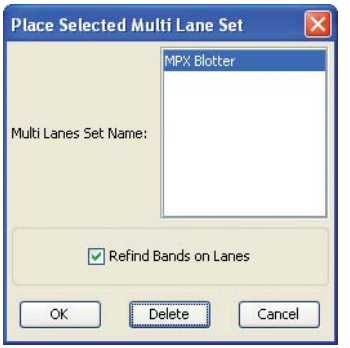
- Create all lanes using the single- or multi-lane creation tools. The multi-lane tool works well for sample lanes from the LI-COR MPX Blotter (or similar blotting systems) since the lanes are uniform.
- Edit the band markers. Add or delete band markers as needed. Adjust band marker size as needed.
- Apply molecular weight standards to the standard lane.

After completing the tasks above, click  or choose **Lane > Save Lanes as Template** to save the current lanes and molecular weight standards to a template.

*Enter a name and click **OK**.*

Placing Lanes Using a Template

To place lanes from a template, click  or choose **Lane > Place Multi Lanes**.



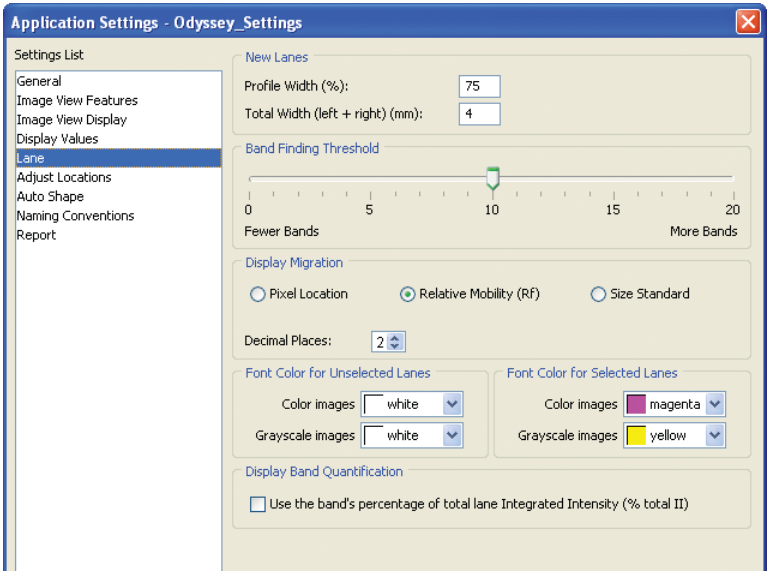
In the template selection window, select a template from the list and click **OK**. Unless **Refind Bands on Lanes** is selected, lanes and band markers are placed in exactly the same position as the original analysis and molecular weight standards are applied in the same way. If **Refind Bands on Lanes** is selected, lanes are placed in the same position, but the stored band markers are discarded and the automatic band finding software is used to find bands within the lanes.

Deleting a Lane Template

Click  to open the template selection window. Select a template and click **Delete**.

Using the Application Settings

To open the Application settings for lanes, choose **Settings > Application** and then select **Lane** from the **Settings List**.



Application settings for lanes are used to control the characteristics of a lane when it is first created. Adjusting the Application settings for lanes can increase band finding accuracy and make the initial lane boundaries more closely match the lanes in the image.

Profile Width

The **Profile Width** field is the percentage of the lane width (starting from the center) that is used to generate the lane profile. The default value of 75 seldom needs to be changed, but unusual band shapes could require a higher or lower percentage in order to generate an accurate lane profile.

Total Width

Each new lane is created with a width, in millimeters, equal to the **Total Width** parameter. Setting **Total Width** to match the typical lane width on your images minimizes the lane width adjustments that need to be made when creating new lanes.

Band Finding Threshold

The **Band Finding Threshold** slider is used to control the band finding software in order to more accurately find the correct number of bands in a lane. If the threshold is set too high, too many bands will be found when new lanes are pasted or created. If the threshold is set too low, too few bands will be found. The proper threshold setting for a given lane can be determined using the Lane Profile window (described earlier in this chapter). After using the profile window to observe typical thresholds for a variety of scans, the default value can be set more accurately.

Display Migration

Band positions on images with bands in lanes can be specified in one of three units: pixel location, percent of lane, or as a molecular weight value (assuming size standards are assigned). The number of decimal places used when the value is displayed can be changed using the **Decimal Places** field. **Pixel Location** is the number of scan lines (rows of pixels) from the top of the image. When **Relative Mobility** is chosen, the top of the lane is 0% and the bottom 100% and bands are assigned a percentage based on position between the top and bottom of the lane.

Note: In the Image View, the band migration value is labeled **Pix Locn** when **Pixel Location** is selected, **Rel Mob** when **Relative Mobility** is selected, and **MW** when **Size Standard** is selected.

Lane Color

The lane border color and font color can be changed for lanes that are selected or unselected using the two sets of font color controls. Controls are provided to pick colors for both color and black-and-white image display styles (some colors do not work well for both display styles). Changing the lane/font color can make it easier to distinguish lanes from band markers. (Choose **Image View Features** in the Application settings list to set band marker color.)

Displaying Band Quantification as a Percentage

When **Use Band's Percentage of Total Lane Integrated Intensity** is selected, the data value for each band is not integrated intensity, but rather the band's percentage of the total integrated intensity of all bands in the lane. The percentage is displayed in the tool tip as the cursor passes over a band and is substituted for the Integrated Intensity quantification value if the display of quantification values (**Settings > Application > Image View Features**) is enabled.

Image View Display Settings for Lanes

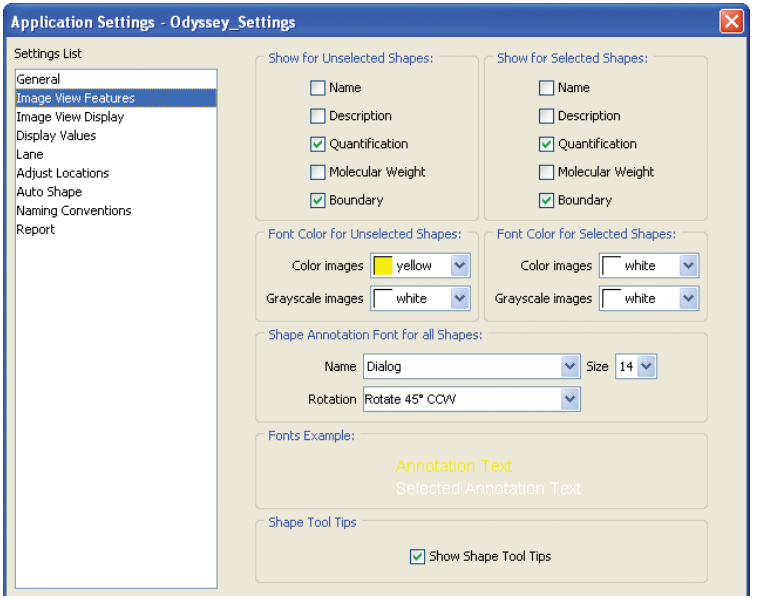
In addition to Lane settings, a useful setting for lanes can be found in the Image View Display settings of the Application settings. If lane names or annotations are truncated at the edge of the image, choose **Settings > Application** and select **Image View Display** from the **Settings List**. Select **Small** or **Large** extended text area around the image (see Chapter 11).

Chapter 6: Band Sizing

Before bands can be sized, an analysis with lanes defined must be open. All bands within the lanes are automatically found when the lanes are created. Creating lanes and finding bands is described in Chapter 5.

Checking the Application Settings

Before sizing bands on an image, it is a good idea to check the Application settings to see how annotations will be displayed. Click  on the toolbar to open the Application settings for image view features.



The settings for image view features can also be opened by choosing **Settings > Application** and then selecting **Image View Features** in the **Settings List**.

The settings for image view features are an important part of any analysis because they can be used to reduce the "screen clutter" that can occur when annotations are displayed on closely spaced screen objects. There are two groups of Image View settings – those for selected screen features and those for unselected features. One strategy to reduce screen clutter is to treat selected and unselected features (i.e. band markers, etc.) differently. If all annotations are turned on for selected features and all annotations except boundaries are turned off for unselected features, the display will stay uncluttered and each individual feature can be clicked to display its annotation (molecular weight, etc).

In other situations, like exporting annotated images, it is useful to have annotations turned on for unselected features. For example, before exporting an image it may be advantageous to turn on molecular weight (MW) values for unselected features so all MW values will be shown on the exported image.

Each of the Image View Feature settings is described below:

- **Name:** Name is either the auto-entered, default object name, or a user-entered name in the object properties (**Edit > Properties**).
- **Description:** Description is blank by default for many objects, but can be entered in the object properties. Descriptions can be included in reports.
- **Quantification:** Quantification values are displayed as Integrated Intensity or Concentration, depending on the Application settings for Display Values. This annotation is generally turned off for band sizing unless bands in the lanes are to be quantified.
- **Molecular Weight:** The Molecular Weight annotation displays band migration in the gel as molecular weight, scan line on the image, or percentage of the lane according to the Application settings for Lanes

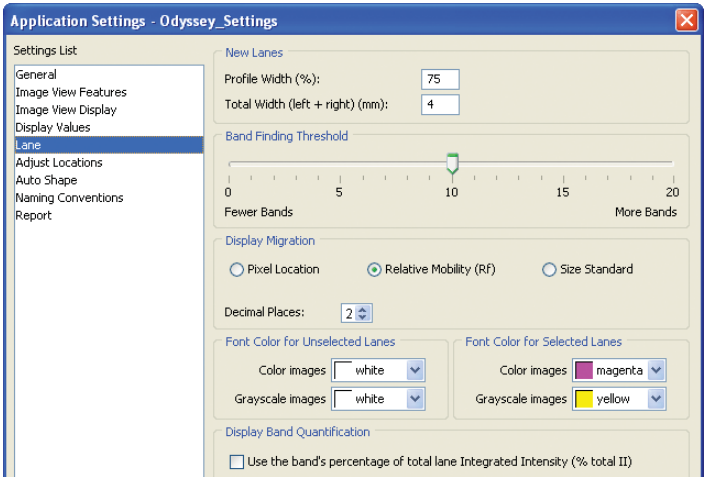
(described below). If migration is displayed as molecular weight, values are displayed as “n/a” (not assigned) until molecular weight standards have been identified.

- **Boundary:** Each feature (i.e. lane, circle, rectangle, band marker, etc.) has a boundary. If boundaries are turned off, only the text annotations will remain.

Turning off the tool tips (**Show Shape Tool Tips**) or changing the font used for annotations can also reduce screen clutter. Changing the normal or selected color of the font, reducing the size of the font, or even changing to a different font can improve the readability of notations on the image. A sample of the currently selected font and font colors is shown in the **Fonts Example** area. Separate controls are provided to pick font colors for both color and black-and-white image display styles (some colors do not work well for both display styles). Annotations can be rotated 45 or 90 degrees counter-clockwise using the **Rotation** list.

Checking the Display Migration Settings

Odyssey can display the migration of a band through the gel as a molecular weight value, scan line (pixel row) on the image, or a percentage of the lane. To change the way band migration is displayed, choose **Settings > Application Settings** and then select **Lane** from the **Settings List**.



Band positions can be specified in one of three units in the Image View, lane profiles, and reports. When **Display Migration** in the **Lane** settings is set to **Pixel Location**, band positions are reported as the number of scan lines from the top of the image. When **Relative Mobility** is selected, the top of the lane is 0% and the bottom 100% and bands are assigned a percentage based on position between the top and bottom of the lane. When **Size Standard** is selected, band positions are reported as a molecular weight value, assuming size standards have been assigned. The number of decimal places used when the value is displayed can be changed using the **Decimal Places** field.

Note: *In the Image View, abbreviations for Pixel Location, Relative Mobility, and Size Standard are "Pix Locn", "Rel Mob", and "MW" respectively.*

Band Sizing in Single Channel Mode

If a scan has two images, the images can be displayed individually in single channel mode or overlaid in a composite image (Chapter 11).

When an analysis with two images is opened for the first time, the images are overlaid so lanes can be added (Chapter 5). After adding lanes, each image must be analyzed separately in single channel mode.

Switching Image Channels

To display a single image, choose **View > Single Channel** or click  on the toolbar. After a single image is displayed, click  , if necessary, to switch between channels until the channel with the molecular weight marker bands is displayed.

Using Size Standard Sets

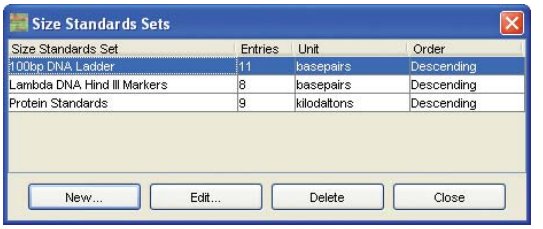
After lanes are defined and bands are edited, the molecular weight (MW) of each MW standard band needs to be assigned. Molecular weights are assigned by adding one MW line for each MW standard band. MW lines, when properly placed, connect all the points of equal MW on the gel. Thus, each intersection between a MW line and the centerline of a lane represents the same MW. For gels where the bands slant or smile, the MW line must be reshaped to follow the contour of the gel as described later in this section.

Note: *Odyssey requires an equal number of MW standard bands in each standards lane. MW sets must also have exactly the same number of standards as there are bands in the standards lanes.*

MW weight lines can be added individually, as described later in this chapter, or in groups that are stored as molecular weight sets. For repetitive scanning with the same molecular weight standards, molecular weight sets save time and eliminate having to enter each molecular weight standard for every image. Creating MW standard lines individually is more useful when a certain set of standards is going to be used for a only few gels.

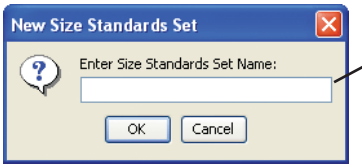
Creating Size Standard Sets

After lanes are created and bands are found, MW sets can be created by choosing **Settings > Size Standard Sets**.



The table in the Size Standards Sets window shows the name, number of standards in the set, units, and sort order for each set.

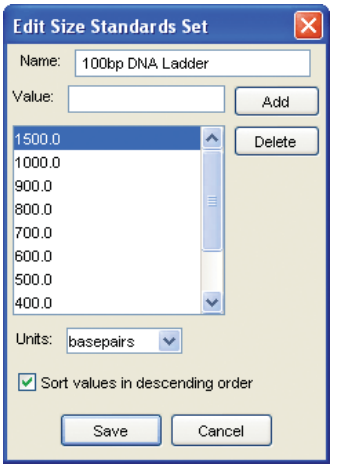
Click **New** to start a new set.



Enter a name for the new MW set and click **OK**.

Names cannot contain any of the following characters: / \ * ? : < > |. It is also not necessary to enter any file name extensions.

The Edit Size Standards Set window can be used to enter the size of all standards in the set, select the units, and specify the sort order for the set.



Begin by selecting the units for the set using the **Units** drop-down list. Basepairs, kilobasepairs, daltons, and kilodaltons can be selected.

Standards are added by entering the value of the molecular weight standard in the **Value** field and clicking **Add**. Standards can be entered as either an integer or a mixed number (integer with a decimal fraction). Continue entering standard values and clicking **Add** until all standards have been entered. The order in which the standards are entered is unimportant since they are automatically sorted.

For typical gels with the highest MW standard at the top of the image and the lowest at the bottom, the **Sort Values in Descending Order** check box should be selected.

If your image has the lowest MW standards at the top of the image, deselect the **Sort Values in Descending Order** check box to reverse the order.

After entering all the standards, click **OK** to save the new set.

Deleting a Standard From a Set

To delete a standard, select the standard in the list and click the **Delete** button.

Editing Size Standard Sets

Any MW standard set can be edited by double-clicking a set in the Size Standard Sets window or by selecting the set and clicking the **Edit** button.)

In the Edit Size Standards Set window standards can be added or deleted as described above under Creating MW Sets.

Deleting Size Standard Sets

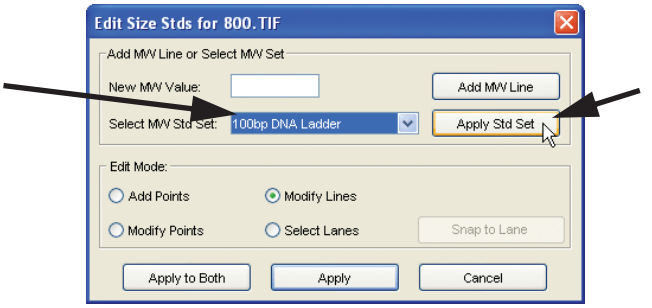
To delete a size standard set, select it in the Size Standard Sets window and click the **Delete** button. If only one standard within a set needs to be deleted, edit the set as described above.

Using Size Standard Sets

Start by choosing **Lane > Edit Size Standards** or by clicking  on the toolbar.

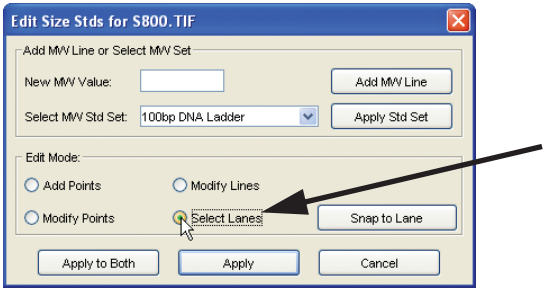
Remember that lanes must be found and only one image displayed before molecular weight bands can be identified.

Choose a MW set from the **Select MW Std Set** drop-down list in the Edit Size Standards window and click the **Apply Std Set** button.



MW lines for all standards in the set are added to the image at once when **Apply Std Set** is clicked. The next step is to identify which lanes contain standard bands and then automatically match the MW lines to the bands in the standards lanes.

To select the lanes containing standards, click **Select Lanes** to set the Edit Mode to **Select Lanes**.



Select <u>all</u> lanes containing size standards. After selecting the first lane, hold down the **Ctrl** key and click additional size standard lanes to select them.

TIP: *Lanes are selected by clicking the centerline, but the centerline is often covered by band markers. Clicking near the top or bottom is usually easier. When the lane is selected, the dashed centerline turns to the highlight color.*

When all the size standard lanes are selected, click **Snap To Lanes**. The MW lines are snapped to the bands in the standards lanes. A control point (square) is added to the center of each MW marker band and the MW line is drawn between the control points.

Note: *In sample lanes, molecular weight standards are assigned at the points where the molecular weight lines cross the centerline of the lane.*

Control points are inflection points on the line that can be moved. For gels with even band migration, no other editing may be necessary since the straight lines between control points accurately represent the location of that molecular weight across the gel. For gels with smiles, the MW lines need to be edited and reshaped so the MW line curves and follows the contour of the gel. Editing MW lines is described below.

Applying Standards to the Image

The MW lines are now shown on the image, but have not been permanently applied. MW lines are applied to only the current image or to both images using the **Apply** or **Apply to Both** buttons (respectively). **Apply** is used when analyzing one image or when each image has its own MW markers. When **Apply to Both** is clicked, the MW lines on the current image are used to size bands on both images (MW bands don't need to be present on the second image).

When **Apply to Both** or **Apply** is clicked, the size standards are identified and all sample bands are automatically sized.

Adding MW Lines One-at-a-Time

The Edit Size Standards window described above can also be used for adding MW lines individually, rather than in sets. For gels with a few size standards that will be run just a few times, it is better to add

size standards individually for each gel, rather than to save a set that would become useless after the gels are run.

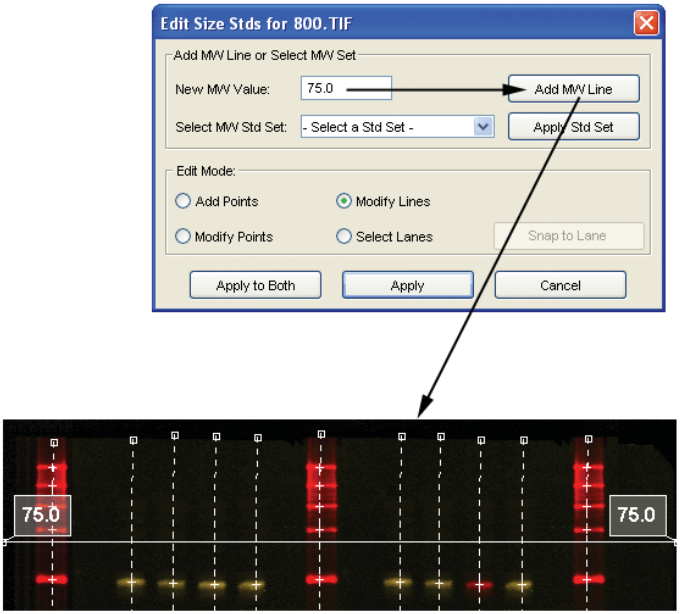
Adding MW lines individually is very similar to using size standard sets, except that each MW line is created individually by entering its MW in the Edit Size Standards window.

As with size standard sets, the number of MW lines added must exactly match the number of bands in the lanes containing standards. After the MW lines are added, the lines are "snapped" to their respective bands and the MW of each sample band is found as soon as the MW lines are applied to the image.

In order to add MW lines, lanes should be found and only one image displayed.

Adding MW Lines

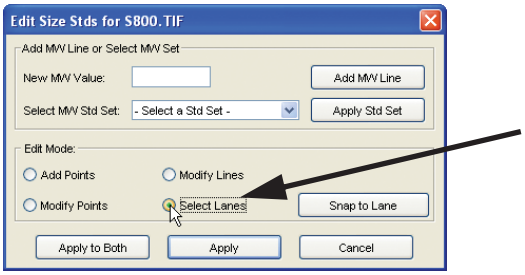
MW lines are added by choosing **Lane > Edit Size Standards**. Individual MW lines are added by entering the size in the **New MW Value** field and clicking **Add MW Line**. Later in the chapter you will see how to change the units in the Size Standards window.



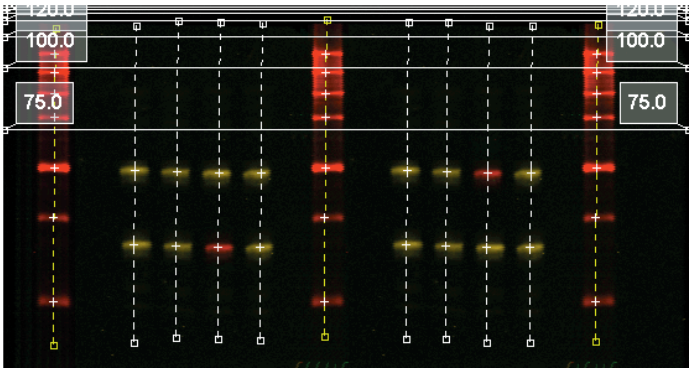
After clicking **Add MW Line**, a MW line is displayed on the image that will connect all the standard bands of a particular weight (75 bp in the example above). The MW line itself represents all the points of equal MW on the gel. The placement and shape of the line is not important yet. After the first line is added, add the rest of the MW lines by entering their MW and click **Add MW Line** as described above.

After all the size lines have been added, the lines need to be linked to specific MW bands in standards lanes. First, the lanes with

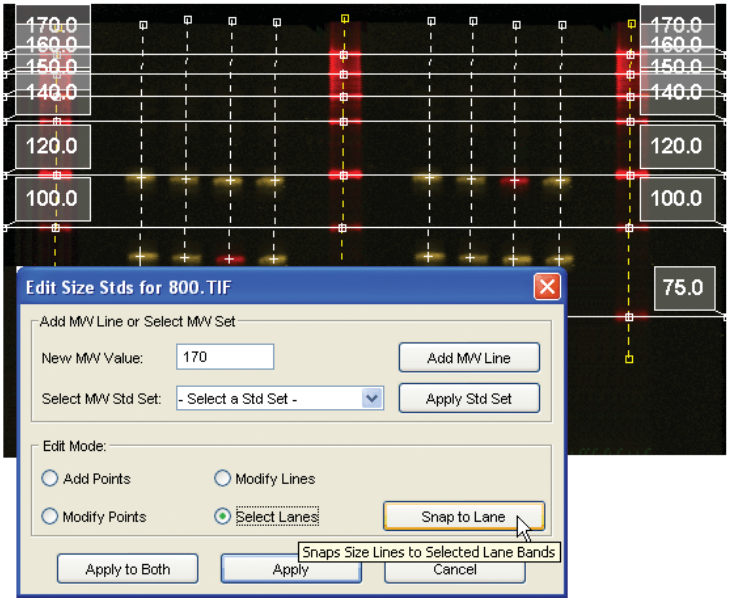
standards need to be selected. Click the **Select Lanes** radio button to set the Edit Mode to **Select Lanes**.



With the edit mode set to **Select Lanes**, the centerline of a lane can be selected by clicking it. To select additional standards lanes, hold down the **Ctrl** key and click the centerline of the lane. All lanes with standards should be selected and there should be no sample lanes selected. If a lane is difficult to select because the center-line is covered with band markers, click near the top or bottom of the lane.



Click the **Snap To Lanes** button in the Edit Size Stds window to link the MW lines to bands in the standards lanes.



When lines are snapped to the bands, control points are added and MW lines are drawn as described above for MW standard sets. After the MW lines are snapped to the bands, the lines must be applied to the image using the **Apply to Both** or **Apply** buttons (see *Applying Standards to the Image* above).

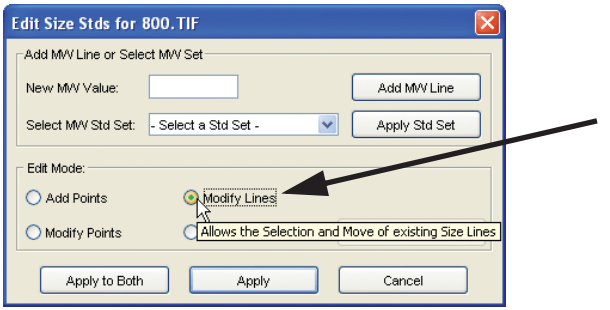
Editing Molecular Weight Lines

On images where fragments had differential migration rates during electrophoresis, bands may "smile", "frown", or be slanted. For band sizing to be accurate, molecular weight lines must follow the contour of the gel, whatever shape that might be. The point at which the MW line crosses the centerline of the lane is assigned the same molecular weight as the line. Therefore, if the MW line crosses the centerline

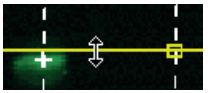
too high or too low, band sizing will be inaccurate. Odyssey software has tools to move and reshape MW lines in the Edit size standards window – the same window used to add the MW lines.

Moving Whole Lines

Whole lines may need to be moved because all points on the line are too high or low. To move a line, set the Edit Mode in the Edit Size Standards window to **Modify Lines** as shown below.



Click a MW line to select it and move the cursor over the line until it becomes an up-down arrow cursor as shown below.



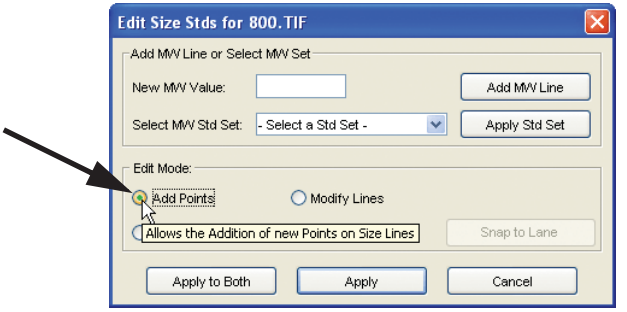
Click and drag the selected line up or down until it is correctly positioned, then release the mouse button. To move more than one line, hold down the **Ctrl** key while clicking multiple lines.

Adding Points to a Line

To accommodate smiles, frowns, or other irregular electrophoresis artifacts, it is necessary to add extra points to the MW line. After the points are added, the next section below shows how to move the

points into new positions that follow the contour of bands on the image.

To add points to MW lines, change the Edit Mode in the Edit Size Standards window to **Add Points**, as shown below.



Click on a MW line at a position where an inflection point needs to be added. Any line can be clicked and any number of points can be added by continuing to point-and-click, as long as the Edit Mode is still set to **Add Points**. To stop adding points, switch to one of the other edit modes like **Modify Points**.

Moving Points

When the Edit Mode in the Edit Size Standards window is changed to **Modify Points**, any individual points can be clicked and dragged to a new position. On images with large smiles or frowns, it may be necessary to add many additional points to make the molecular weight line conform to the curve. (Remember that the point where the MW line crosses the centerline of a lane is where that molecular weight standard is located in that particular lane). For images with smiles or other curves, plotting the size standards (described below)

for each lane will identify any MW standards that are out of position due to misplaced points or lack of points.

Plotting Size Standards

Band sizing is not complete until the Size Standards window has been used to set the size interpolation method and to check the standards plot for each lane. The rest of this chapter discusses how to accomplish these important goals.

Setting the Interpolation Method

When molecular weight standards are applied to an image, sample bands of unknown MW weight are automatically sized using a mathematical method to interpolate the size of sample bands that lie between standard bands. The interpolation method should always be verified by choosing **Lane > Size Standards**.

Selecting the correct interpolation method is important because it affects the size assigned to sample bands. **Linear**, **Log**, **Reciprocal Fit**,

and **Exponential** interpolation methods can be chosen from the **Interpolation Method** list.

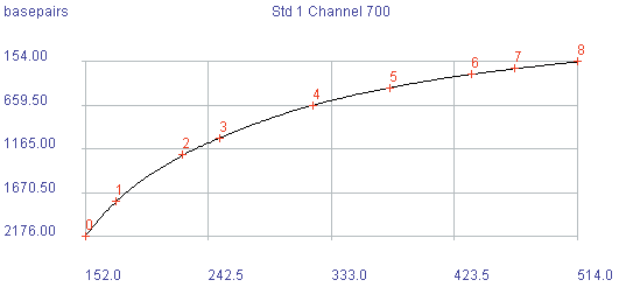
In general, the **Reciprocal Fit** interpolation method usually produces the best results. There are exceptions, however. If the plot of the standards is very linear, the **Linear** interpolation may yield slightly better results. If the image has sample bands with a size that is higher or lower than the standards, use the **Exponential** interpolation method. Any time there is a need to extrapolate beyond the size standards, the **Exponential** interpolation method usually produces the best results. The interpolation applies to all lanes on the image.

Setting Units for Standards

Like the interpolation method, the **Units** apply to all lanes and should be set before examining the plot of the standards for each lane. Basepairs, kilobasepairs, daltons, and kilodaltons can be selected for units.

Reviewing the Standards Plot for Each Lane

The **Lane** drop-down list in the Size Standard plot window is used to select the lane to plot. Start with the lanes containing MW standards. Look for anomalous bands. For example, if the smooth curve in the plot below is broken by a standard that is too high or too low, use the Edit Size Standards window to review the position of the MW line of the lane in question. The line may be snapped to the wrong band or the size may be incorrectly assigned.



After reviewing the plots for lanes with MW standards, review the sample lanes to make sure each standard is correctly positioned. In sample lanes, molecular weight standards are assigned at the points where the MW lines cross the centerline of the lane. For gels with even migration (straight molecular weight lines), the plots of sample lanes will be very similar to standards lanes. For gels with smiles or other gel artifacts where extra points are added to bend the MW lines in a particular direction, close examination of each sample lane is needed.

Suppose you have an image with a smile and too few points are added to the molecular weight line. In this case, the MW line may not accurately follow the contour of the gel and may cross the centerline of a lane at a point that is too high. This will be revealed in the standards plot for the lane in question since the standard in question will be misplaced on the plot. Adding another point to the MW line and moving the point to the proper position will eliminate the problem.

When all standards have been examined, click **OK** to close the Size Standards window. Sample bands already have a MW assigned, so after any adjustments are make to the MW lines, sizing is complete and the MW assigned to each sample band is final.

Chapter 7: Drawing Features on Images

Overview

For images that do not have bands in lanes, analysis begins by drawing individual features (circles, squares, etc.) that surround all the fluorescent dots or bands in the image. Features can be drawn on the image using a shape tool (rectangle, circle, oval, free form shape, etc.). A grid tool is also available to quickly apply features to an image in a regularly spaced grid pattern. The grid tool is ideal for scans of microplates. Each grid can also have a subgrid in each grid cell, making the grid tool useful for protein arrays.

Drawing Features on the Image

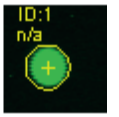
For quantification, each image should be analyzed separately. When adding individual features to an image, only one image should be displayed. Begin by clicking  on the toolbar until only one image is displayed in Single Channel mode (assuming the scan has two images).

Click the shape tool in the toolbar that most closely matches the dot or band on the image. Rectangle  , Circle  , Oval  and Freeform tools are available.

Draw a feature that encloses the dot/band. Features are drawn the same way they are in most drawing programs. Rectangles, circles, and ovals are drawn as shown below.

Imagine a bounding rectangle surrounding the band/dot and place the cursor in the upper left corner.

Click and hold down the mouse button. Drag downward and to the right. Release the mouse button when the fluorescence on the image is completely enclosed.



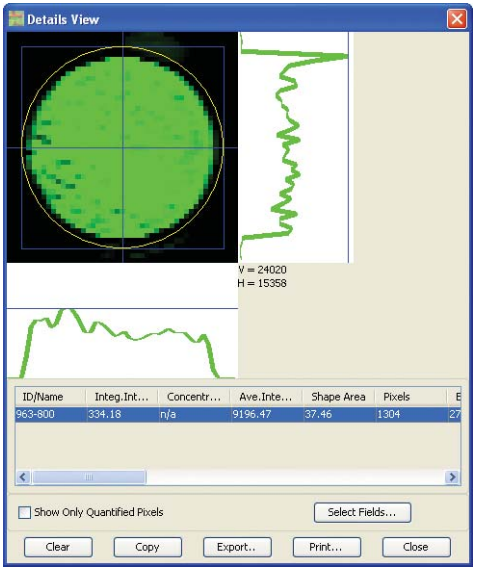
Drawing a feature with the Freeform tool is slightly different. To draw a feature with the Freeform tool, click the Freeform button in the toolbar and move the cursor to the edge of the object in the image. Click and hold down the left mouse button and trace around the object on the image. Release the mouse button when the freeform line completely encloses the object in the image.

Keyboard Shortcut: *The last tool selected from the toolbar is automatically selected again when F5 is pressed.*

Using Details View to Position Features

Odyssey has a Details View that is useful for verifying that a feature is correctly positioned and is large enough to enclose the fluorescence on the image. To open the Details View, select a feature and click  on the toolbar or choose **Report > Details View**. Details View can also be opened by selecting a feature, right clicking the image, and choosing **Details View** from the pop-up menu.

It is advantageous to leave Details View open while drawing features on an image because each new feature is immediately shown in Details View after it is drawn.



Details View shows an enlarged view of the dot/band along with the shape that surrounds it. The center of the feature surrounding the dot/band is marked by the blue cross hairs. The sides of the blue rectangle are the horizontal and vertical boundaries of the feature on the image. In the case of a rectangle feature, the blue line actually represents the rectangle. The curves to the right and below the image plot the intensity of the pixels below the cross hairs.

First, make sure the feature on the image is large enough to enclose all the fluorescence. To get the best background calculation for quantification, all pixels on the outside perimeter of the blue line should be empty background pixels (no fluorescence). For smeared bands or closely spaced dots this may not be possible, but the background calculation method can be changed to accommodate the image data, as described in the Chapter 8.

After the feature is properly sized, examine the cross hairs to make sure they are centered over the band/dot. If a feature needs to be

moved to center it in the cross hairs, the feature must be moved on the image rather than in Details View.

Resizing and Deleting Features

If a feature (circle, etc.) is too large or too small, move the cursor toward a corner until the cursor turns to a diagonal arrow as shown below. With the diagonal arrow cursor displayed, the feature can be enlarged or reduced by clicking and dragging.

*To delete a feature, select the feature and press the **Delete** key on the keyboard or click* ✕ *on the toolbar.*

Moving Features

If the cross hairs in Details View indicate that a feature needs to be moved in order to center it, move the cursor to the center of the feature on the image until the cursor has arrows in all four directions, as shown below. When the "all-arrows" cursor is displayed, features can be clicked and dragged to a new position.

Nudging Features: It can be difficult to precisely place objects with the mouse. Selected features can be nudged one pixel at a time using the arrow keys on the keyboard, as long the all arrows cursor is displayed within the border of the feature.

Selecting Multiple Features: Multiple features can be selected by clicking and dragging a selection rectangle around them, or by control clicking additional features after selecting the first feature.

Automatically Adjusting Feature Locations: Although features can be moved manually, it is often easier let Odyssey software adjust the locations of selected features. As described later in this chapter, choosing **Analyze > Adjust Feature Location** automatically moves features over any nearby fluorescence in the image. Successful use of the automatic adjustment software depends on the features being near the fluorescence that should be enclosed and large enough to enclose the fluorescence without being so large that extraneous fluorescence gets enclosed.

Copying and Pasting Features

With one exception, selected features can be copied and pasted, or cut and pasted, in the normal way. The exception is the location at which the feature is pasted. Features are pasted with the upper left corner of the feature at the current position of the mouse cursor. The normal paste procedure is to move the mouse cursor to the desired position and press **Ctrl+v** on the keyboard.

A variation of the **Paste** command is the **Paste Special** command. **Paste Special** pastes features in exactly the same pixel locations as the original features. **Paste Special** is useful for pasting features between images. For example, if one image has an array of dots with circles drawn circles around them, the circles can be copied to the second image by doing a **Select All**, copying the circles (**Ctrl+c)**, switching images and doing a **Paste Special**. All the circle features will be pasted in precisely the same location as the first image, eliminating the need to draw all the circle features on the second image.

Multiple Feature Selection: When multiple features are selected, they can be copied and pasted as a group. To select multiple features:

- Click the first feature. Hold down the **Ctrl** key and click additional features.
- Click and drag a selection box around all the features to be selected.

After copying and pasting features, they can be automatically moved into the correct position using the **Adjust Feature Location** function described later in this chapter.

Adding Multiple Features

Setting the Drawing Mode to Continuous

The drawing mode tool ( ) can be used to toggle the drawing mode for features (shapes) between drawing single features and drawing a new feature each time you click and drag across the image. To determine which mode is currently selected, move the cursor over the drawing mode tool and read the tool tip. A menu choice on the **Analyze** menu also toggles between **Single Shapes** (indicating continuous mode is selected) and **Continuous Shapes**.

Suppose you want to draw five circles and the mode is set to Single Shapes. Begin by clicking the drawing mode tool ( ) to change to Continuous Shapes, click the circle tool, and then click and drag on the image to create the first circle. After releasing the mouse button, click and drag again for the second shape and repeat until all five shapes are drawn. Click to resume drawing single shapes.

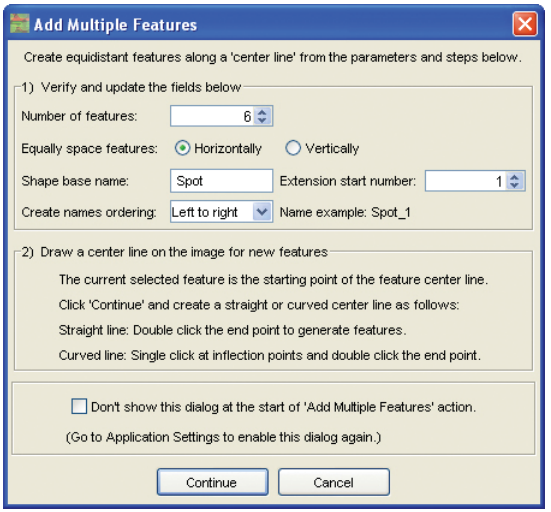
Note: *The drawing mode tool works with only the circle, ellipse, rectangle, freeform and text tools.*

Using the Multiple Features Tool

When there are equidistant image features (bands, spots, etc.) along an imaginary straight or curved line, the multiple features tool ( ) can be used to add a user-specified number of equally spaced features (circles, etc.).

Features are added along a straight or curved line that starts from the center of an existing feature and ends at a point where the user double clicks the mouse. The first step is to add the first feature and select it. Since this feature is one end point of the imaginary line connecting the image features, it should be on the left, right, top, or bottom side of the line.

After selecting the first feature, click the multiple features tool ( ) or choose **Analyze > Add Multiple Features**.



In the Add Multiple Features window, start by specifying the number of features to add and whether the features should be spaced equally in the horizontal or vertical dimension. Features are named with a prefix (base name) and a numbered extension (see **Name Example** in the window above). Name extension numbers are incremented for each feature, moving in the specified direction: left-to-right, right-to-left, top-to-bottom, or bottom-to-top.

Adding Features Along a Straight Line

The next step is to draw the line. Click **Continue** to return to the image. For a straight line, double click in the center of the spot/band

at the opposite end of the line from the initial feature. The imaginary line connecting the two points can be horizontal, vertical or any angle in between.

Adding Features Along a Curved Line

For curved lines, click **Continue** and then single click at inflection points along the curve until reaching the spot/band on the end opposite the initial feature. Double click at the end point to exit line draw mode and the specified number of features will be drawn.

Disabling the Add Multiple Features Window

For applications with repetitive sample configurations, the Add Multiple Features window may not be necessary and can be disabled by selecting **Don't show this dialog at the start of 'Add Multiple Features' action**. For example, if many blots with multiple rows from an 8-channel pipettor are scanned, there is no reason to specify to add seven features for every row. With the Add Multiple Features window turned off, all that is necessary is to add the initial feature, click the Multiple Features tool, and double click in the center of the last spot. The latest configuration (number of features, names, etc.) of the Multiple Features tool is always stored in memory for the next use.

Note: *If the Add Multiple Features window is turned off, it can be turned back on by choosing* ***Settings > Application****, selecting* ***General*** *from the* ***Settings List****, and deselecting the* ***Don't show the Create Multi Features Dialog*** *check box.*

Automatically Adjusting Feature Locations

When placing features (circles, squares, polygons, etc.) on an image, it is not necessary to place them precisely because selected features can be moved to enclose nearby fluorescence by choosing **Analyze > Adjust Feature Location**. After selecting the features, **Adjust Feature**

Location can also be invoked by clicking  on the toolbar or right clicking the image and choosing **Adjust Feature Location** from the pop-up menu.

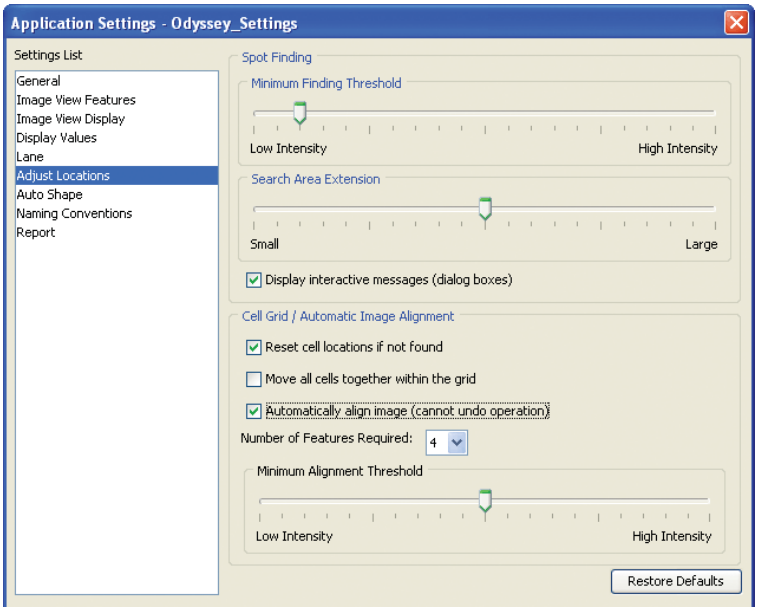
Note: *The* ***Adjust Feature Location*** *function moves features only. It does not size them. If the initial size is too small to enclose the fluorescence, features will not be moved to optimal locations.*

Tip: *If features are misplaced, choose* ***Edit > Undo****.*

Note: *Even though* ***Adjust Feature Location*** *can be used with grids or features within grids, it generally does not place features accurately due to the fluorescence of the plate walls.*

Using the Adjust Location Settings

The locations of selected features on the image are moved based on a localized search for maximum signal. The Adjust Locations settings are used to refine the search criteria and improve spot finding. To open the Adjust Location settings, choose **Settings > Application** and select **Adjust Location** from the **Settings List**.



Improving Spot Finding

The Adjust Location settings determine the search area (how far from the original location to search) and threshold intensity for spot finding.

The **Minimum Finding Threshold** slider sets the threshold intensity for spot finding. Since images vary in background and fluorescent intensity, this setting must be determined empirically for your images. Spurious fluorescence due to dirt, fluorescence of microplate walls, and other fluorescent sources needs special consideration. For example, dirt that fluoresces brightly near a selected feature can cause the feature to be moved to the wrong location. Setting the threshold to a higher intensity may prevent features from being placed over spurious fluorescence, but if the threshold is set too high, features near low intensity fluorescent dots may be

misplaced. Feature locations may need to be adjusted manually on images with a lot of bright spurious fluorescence near image features.

Keeping the search area small is generally desirable, even though it means the initial feature locations must be near the final locations. If the search area is too large, the search may find a bright piece of dirt some distance away from the target fluorescence.

The **Search Area Extension** slider is used to increase or decrease the image area around a selected feature that is searched for maximum fluorescent signal. The area searched is defined differently for features within grids and features drawn on the image. For a feature within a grid, setting **Search Area Extension** to **Small** confines the search area to the boundaries of the grid cell. Setting **Search Area Extension** to **Large** expands the search area to twice the height and width of one grid cell, centered in the center of the original grid cell. For a feature drawn on the image, setting **Search Area Extension** to **Small** extends the search in all directions (top, left, bottom, right) by one quarter of the minimum dimension of the feature. Setting **Search Area Extension** to **Large** expands the search in all directions by the minimum feature dimension. Enlarging the search area slows the software.

Note: *If Adjust Feature Location is used with microplate images that have fluorescence from the microplate walls, grid features may not be accurately placed no matter how the threshold and search area are configured.*

While new feature locations are being calculated, a variety of warnings or error messages may be displayed depending on the image features and original location of the features. To suppress these messages, deselect **Display Interactive Messages**. (Messages can still be viewed by double clicking the message bar at the bottom of the Odyssey Window to display the Status Message History window.)

Grid Features and Image Alignment

For grids, the settings can also be used to move all the selected grid features by an equal amount and to reset features to their original location if not found.

If **Reset Cell Locations If Not Found** is selected, selected features in a grid are moved to the center of their respective grid cells if a location cannot be found using the **Minimum Finding Threshold**. Features not associated with grids are never moved if their location cannot be found.

All features within grids are moved the same distance when **Move All Cells Together Within The Grid** is selected and a consistency threshold is met. Features will not be moved unless a user-defined number of features are found to require approximately the same movements for correct positioning. The number of features that must exhibit consistent movement is entered in the **Number of Features Required** field.

When the location of features in a grid are adjusted, images are automatically aligned if **Automatically Align Images** is selected and the number of features requiring the same offset is equal to or greater than the number of features specified in the **Number of Features Required** field.

After alignment, the fluorescent features of the two images should be in the same positions. The status line at the bottom of the Odyssey window indicates which image was offset and by how much, assuming the alignment was successful (in some cases it may be necessary to double click the status line to open the message history window).

Important: There is no undo operation for the image alignment routine because the TIFF image files are immediately changed.

Restoring Default Adjust Location Settings

Click **Restore Defaults** to change the Adjust Location settings to the values shown in the Adjust Locations window at the beginning of this section.

Adding Multiple Features Using Grids

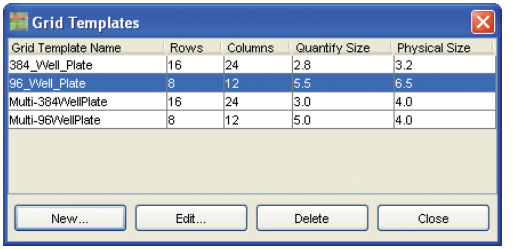
Odyssey's grid tool can be used to apply an array of features, all at once. For regularly spaced image features, such as those created by microplates or manually placed spots in dot blots, the grid tool can save considerable time compared to manually placing each feature using the shape tools. For protein arrays, the grid tool can also be used to apply subgrids, which are grids within each cell of a larger grid.

Grids are often applied to images for In-Cell Western assays or other analyses that have independent background calculation methods. To accommodate this, the background calculation method (**Analyze > Background Method**) is changed to **No Background** when a grid is added (either manually or automatically as described below). If you want to use one of the other background calculation methods, apply the grid, select the background shape, and set the background method by choosing **Analyze > Background Method**.

Before a grid can be applied, a grid template must be defined that specifies the size and placement of the grid, as well as the size of the features and whether the features are circles or rectangles. Grid templates for typical 96- and 384-well plates are supplied with Odyssey software and can be modified as needed to create new templates.

Creating Grid Templates

The Grid Template settings are used to save grid templates. The Grid Template settings are opened by choosing **Settings > Grid Templates**.



A grid template can be created using one of the following methods:

- Copy an existing template. Select a template in the Grid Templates window, click **Edit**, change the grid parameters (listed below), and click **Save As** to name the edited template.
- Click **New** in the Grid Templates window, change the grid parameters, and click **Save As** to name the new template.

Grid templates can also be saved from the main Odyssey window as follows:

- After placing a grid and adjusting it to match the image, leave the grid selected and open the grid properties to save the grid parameters to a grid template. Grid properties are opened by right clicking the image and choosing **Properties** or by clicking the properties button ( ) on the left toolbar. After opening the properties window, click **Create Template** to open the **Modify Grid Settings** window with all the parameters filled in to match the grid selected in the Image View window. Review the parameters, click **Save As**, and name the new template.
- A special type of grid template can be saved by selecting a grid, right clicking the image in the Image View window, and selecting **Save Grid As Template** in the popup menu (or choose **Analyze > Save Grid As Template**). This method can be used to visually adjust

the grid and save the grid settings after the grid is correctly positioned. This method also stores additional grid parameters that cannot be entered in the Grid Template settings. For example, if you rotate a grid, right click the image, and choose **Save Grid As Template**, the amount of rotation will be stored in the template, even though rotation is not a parameter that can be entered in the Grid Template settings. Any changes become part of the template when **Save Grid As Template** is clicked.

Note: *Templates saved using* ***Save Grid As Template*** *are not editable; nor can they be used for creating subgrids.*

Deleting Grid Templates

To delete a template, select the template in the Grid Templates window and click **Delete**.

Editing a Grid Template

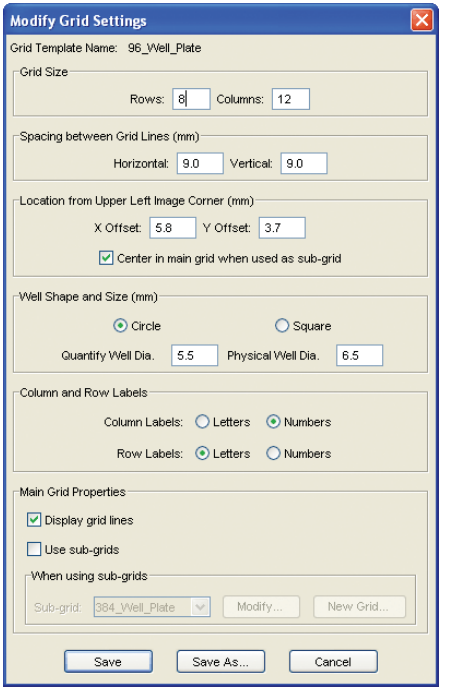
1) Select a template from the template list in the Grid Templates window and click **Edit**.

2) Change the grid parameters as needed.

3) Click **OK** to save the changed grid template.

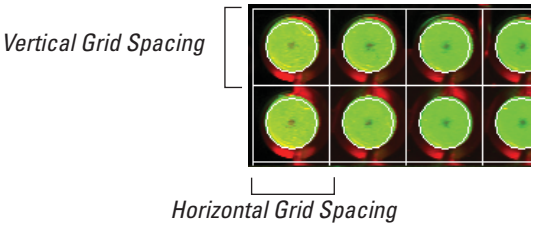
Grid Parameters

When you edit or create a grid template, the grid parameters are listed in the Modify Grid settings window. Each of the parameters are discussed below.

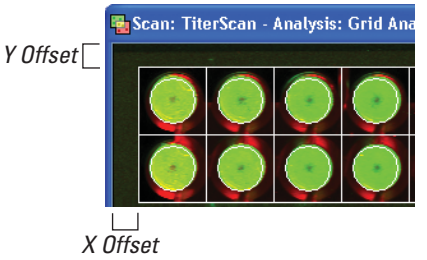


Grid Size: Enter the number of rows and columns in the array of image features. For example, to create a template for a 96-well microplate, you would enter 8 rows and 12 columns (horizontal orientation).

Grid Spacing: The vertical and horizontal spacing (in mm) of the grid lines can be changed in order to match the spacing between objects on the image. The vertical and horizontal spacing should be the same as the vertical and horizontal center-to-center distance between objects on the image. See *Measuring Size and Distance on the Image* below.



Offset From Upper Left Image Corner: These X and Y coordinates determine the placement of the upper left corner of the grid. The offset, in millimeters, is measured from the upper left corner of the image.



Well Shape and Size: Either circle or square features will be placed in the center of each cell on the grid depending on whether **Circle** or **Square** is selected for the well shape. The physical size of the well/spot is the actual size (mm) of the feature to be added to each grid cell (normally just large enough to surround all the fluorescence). Physical size is used to accurately place the grid. The quantification size specifies the size analyzed during quantification and is often the same as the physical size. However, if there is background fluorescence from the microplate in the well edges, setting the quantification size slightly less than the physical size will eliminate quantification of background signal. See *Measuring Size and Distance on the Image* below for details on measuring the size of image features.

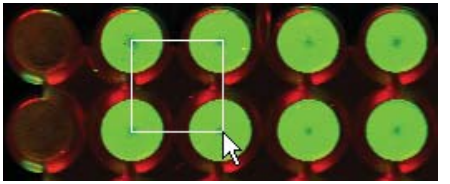
Column and Row Labels: These buttons control whether the row and column labels on the outside of the grid are numbers or letters.

Display Grid Lines: When this check box is selected, grid lines are shown on the image.

Use Sub-grids: The **Use Sub-grids** check box and the **Sub-grid** drop down list for specifying a sub-grid are discussed below in *Using Sub-grids*.

Measuring Size and Distance on the Image

Size and distance values in the grid settings must be fairly precise for optimal operation, but these values are difficult to visually estimate. Size and distance on the image can be determined precisely using the selection rectangle.



Example for measuring grid spacing.

To measure grid spacing, offset, or feature size, drag a selection rectangle over the area to be measured. The X-Y dimensions of the selection rectangle will be displayed in the status message at the bottom of the Odyssey window as shown below.

The measured values give a good starting point that can be refined by applying a grid to the image and adjusting the grid template as needed.

Applying Grids to Images

Grids are always applied to both images in exactly the same location, so it is best to have both images overlaid before applying the grid (use  in the toolbar). After defining a grid template, a grid can be applied to an image by clicking the grid tool () on the toolbar or choose **Analyze > Add Grid**.

In the Select Grid window, select the grid template from the **Grid Name** drop-down list and click **OK** to apply the grid to the image.

To change the grid template before applying it, click the **Modify** button in the Select Grid window and change the grid parameters as described above under *Grid Parameters*.

After clicking **OK**, the grid is automatically placed on the image with the upper left corner at the X,Y offset specified in the grid template.

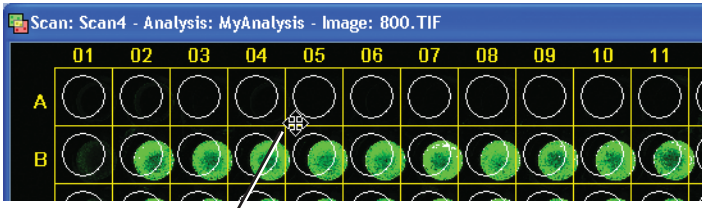
Applying a Grid Automatically

If you have the In-Cell Western Module for Odyssey Software, grids can be applied automatically (Odyssey software finds the correct position) as described in Chapter 9.

Moving a Grid Manually

If the grid is not placed in an optimal position, start by clicking the grid to select it. Next, move the cursor inside the grid so the all-arrows cursor is displayed. With the all-arrows cursor displayed, click and drag the grid until the circle or rectangle features are in the best possible alignment with the objects in the image.

Tip: *While the all-arrows cursor is displayed, the grid can be nudged one pixel at a time using the arrow keys on the keyboard.*



All arrows cursor.

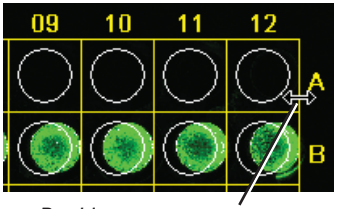
Note: *It is not necessary for features on the image to be centered in grid cells. The grid is provided merely as a visual aid and is not used in analysis. The only thing that is important is that the circles or squares in the grid surround the fluorescence in the image.*

Deleting a Grid

To delete a grid, select it and click the delete button (✕) on the tool bar. Pressing the **Delete** key on the keyboard also deletes a selected grid.

Resizing a Grid

To reshape a grid, first select the grid by clicking one of the grid lines. Move the cursor over one of the border lines of the grid until the double-arrow cursor is displayed.



Double-arrow cursor.

With the double arrow cursor displayed, click and drag the left or right border line to resize the grid horizontally. Click and drag the upper or lower border line to resize the grid vertically.

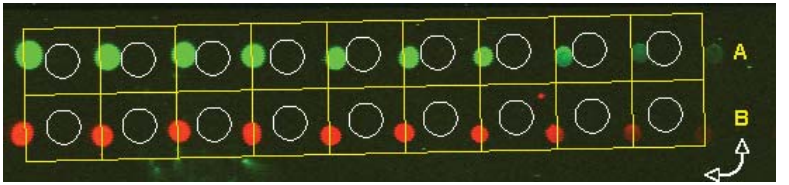
Note: *A grid can also be resized by deleting it, modifying the grid template and applying the grid again.*

Rotating a Grid

To Rotate a grid, select the grid and move the cursor over the lower right corner of the grid until the curved rotation cursor is displayed.

Rotation cursor.

With the rotation cursor displayed, click and drag the cursor upward or downward to rotate the grid.



Moving Features

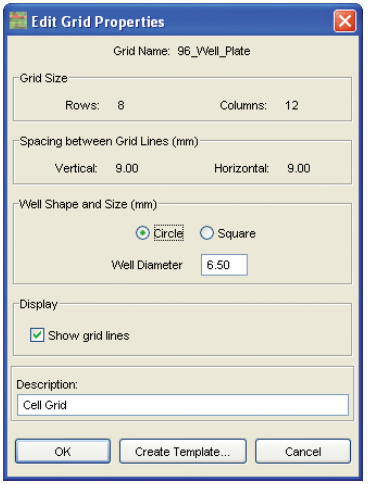
To move individual features, first click the feature to select it and move the cursor inside the feature until the all-arrows cursor is displayed. With the all-arrows cursor displayed, click and drag the feature until it encloses all the fluorescence on the image.

*Multiple features can be selected by holding down the **Ctrl** key while clicking additional features, or by dragging a selection rectangle around multiple features.*

Note: *The circle or rectangle features in the grid do not have to stay within the boundaries of the grid cells. The only thing that is important is that the features fully enclose the fluorescence in the image. The grid lines are only a visual placement aid.*

Changing the Feature Size or Type

The size of a feature or type of feature can be changed after a grid is applied using the grid properties. To open the grid properties, select the grid and click the properties ( ) button. Right clicking the image and choosing **Properties** from the popup menu also opens the grid properties.



The grid size and spacing are listed in the Properties for information only. To change the number of rows or columns in a grid, delete the grid, change the Grid Template settings, and apply the grid again. Grid spacing can be changed by reshaping the grid as described above in *Resizing A Grid*.

If the initial features are too large or small when the grid is applied, the **Well Shape** and **Size** fields can be used to change the size or even

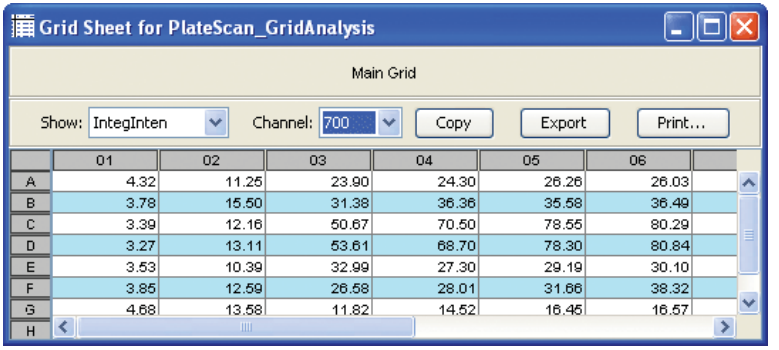
the type of feature. To change from circles to squares, or vice versa, select **Circle** or **Square**. To change the size of the feature, enter the **Well Diameter** in millimeters for circles or the **Well Width** and **Height** for squares. The grid description can be changed in the **Description** field.

IMPORTANT: When **OK** is clicked to apply the properties, a new grid is applied that uses the new features and well sizes. Any changes to the existing grid will be lost.

Displaying Grid Data in the Grid Sheet

Grid objects are often closely spaced, making it difficult to display data values next to the features on the image. For this reason, data are not displayed on the image. Instead, a *grid sheet* is used to display data in table format for each feature in the grid. A grid sheet can be opened by selecting the grid and then choosing **Analyze > GridSheet** or the grid sheet button () on the toolbar.

The numbering or lettering of rows and columns in the grid sheet corresponds to rows and columns in the grid.



	01	02	03	04	05	06
A	4.32	11.25	23.90	24.30	26.26	26.03
B	3.78	15.50	31.38	36.36	35.58	36.49
C	3.39	12.16	50.67	70.50	78.55	80.29
D	3.27	13.11	53.61	68.70	78.30	80.84
E	3.53	10.39	32.99	27.30	29.19	30.10
F	3.85	12.59	26.58	28.01	31.66	38.32
G	4.68	13.58	11.82	14.52	16.45	16.57
H						

Data for one channel at a time is displayed in the Grid Sheet. To switch between channels, use the **Channel** drop-down menu. The **Show** menu can be used to display integrated intensity, concen-

tration, or raw intensity values (see Chapter 13 for calculation descriptions).

The values shown in the Grid Sheet can easily be transferred to a spreadsheet for plotting or analysis. Click **Export** to create a text file with tab-separated values. The text file will contain all data in the Grid Sheet, plus a header that describes the scan. Click **Copy** to copy all data to the clipboard. Individual rows can be selected by Control-clicking the row(s) and pressing **Ctrl+c** to copy the data to the clipboard. Reports and the "Microsoft Excel for Grids" plug-in report can also be used to export grid data. The **Print** button sends the entire grid sheet to the specified printer in the page orientation of your choosing.

Changing Font Size in the Grid Sheet

The font size used in the Grid Sheet can be changed by choosing **Settings > Application** and changing the **Grid Sheet Font Size** setting in the **General** settings.

Copying Grids Between Analyses

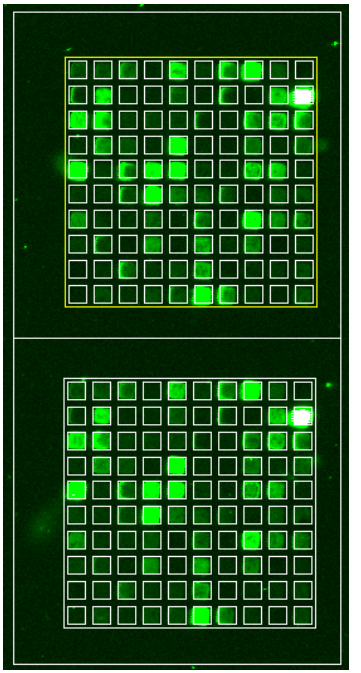
The easiest way to copy a grid from one analysis to the other is to select the grid, choose **Edit > Copy**, switch to another analysis and choose **Edit > Paste** to paste the grid into the second analysis.

IMPORTANT: Any existing grid in an analysis will be replaced when a new grid is pasted.

Grids can also be saved for use in future analyses by selecting the grid and choosing **Analyze > Save Grid as Template**. The new grid template can then be applied using the grid tool like any other grid template. Note, however, that templates saved in this fashion cannot be edited.

Using Subgrids

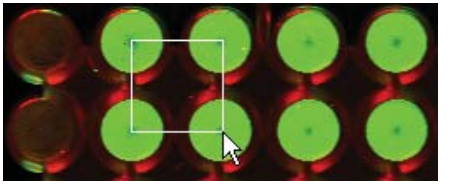
In the image to the right, there is one main grid (2 rows x 1 column) and each cell in the main grid contains a subgrid (10 rows x 10 columns). A strategy for creating subgrids is outlined below.



Designing a Subgrid

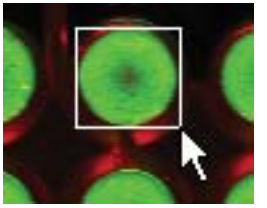
A subgrid cannot be resized after it is included in a main grid, so the first task is to create the subgrid and adjust its size.

1) Click and drag a selection rectangle as shown below to measure the grid spacing. The width and height of the rectangle will be listed in the status message at the bottom of the Odyssey window.



Example for measuring grid spacing.

2) Measure the size of a well/spot by dragging a selection rectangle that encloses the fluorescence as shown below.



3) Choose **Settings > Grid Template** and click **New** in the Grid Templates Window to create a new template for the subgrid.

4) Enter the number of rows and columns in the subgrid.

5) Enter the horizontal and vertical spacing measured in Step 1.

6) Select the type of feature (circle/square) and enter the value measured in Step 2 for the "physical" and "quantify" size (vary the “quantify” size if you need to quantify an area that is less than the entire well/spot).

7) Enter a small number, such as 0.5, for X and Y offsets. When the subgrid is applied, it will be in the upper left corner of the image, which is fine for this design phase.

8) Select the **Center in Main Grid When Used as Sub-grid** check box so the subgrid will be centered in the cells of the main grid when the main grid is applied.

9) Click **Save** and name the template for the subgrid.

10) In the Odyssey window, click the grid tool and apply the subgrid template to the image.

11) Move the subgrid into position and size the entire subgrid to fit the image as needed.

12) Check to make sure the feature size in the subgrid is correct. If not, click  on the toolbar to open the properties and change the feature size to fit. Repeat as necessary.

13) After the grid and grid features are correctly sized, click  on the toolbar to open the properties and click **Create Template**.

14) Check to make sure the **Center in Main Grid When Used as Subgrid** check box is selected and enter small numbers (e.g. 0.5) in the X- and Y-offset fields. Click **Save As**, enter a name for the new template, and click **OK**.

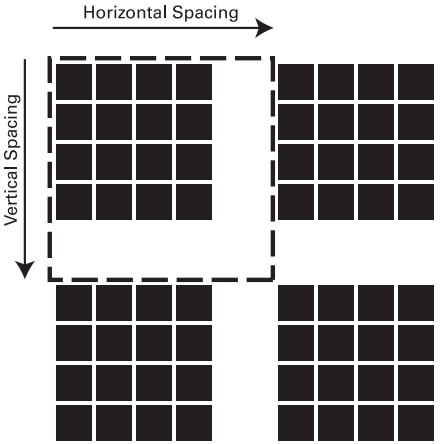
The original subgrid template can either be deleted, or changed and used for the main grid.

Designing a Main Grid

The main grid is specified in a different template than the subgrid. Each cell of the main grid will have a subgrid inside it, so the number of cells in the main grid should match the number of arrays on the image.

Start by measuring the spacing and location dimensions using the selection rectangle as described earlier in this chapter.

1) Measure the horizontal and vertical grid spacing as shown below. Click and drag a selection rectangle starting in the upper left corner of one array. Drag downward and to the right until the next arrays (if any) are contacted as shown in the diagram below.



2) Choose **Settings > Grid Template** and click **New** in the Grid Templates window to create a new template for the main grid.

3) Enter the number of rows and columns (2 x 2 in our example above).

4) Enter the measured dimensions (mm) for horizontal and vertical spacing.

5) Set the X and Y offsets to a small value so the grid is placed in the upper left corner of the display.

6) Select **Use Sub-grids** to add a subgrid in every cell of the main grid and choose the subgrid template from the **Sub-grids** list.

7) Click **Save** and name the template for the main grid.

8) Apply the main grid to the image using the grid tool in the Odyssey window.

9) Move the grid from its initial location and place it so the features in the subgrid are aligned as well as possible with the fluorescence on the image.

Tip: *When a subgrid is selected, right clicking the image opens a pop-up menu that has several useful choices for selecting all or part of a subgrid (single rows, columns, etc.).*

10) Select **Analyze > Adjust Feature Location** to move the features to their final location. If spot finding is not very accurate, select **Edit > Undo Adjust Feature Locations**, change the spot finding parameters in the Application settings (discussed earlier in this chapter), and select **Analyze > Adjust Feature Location** again. On images with a lot of extraneous fluorescence (e.g. microplates with fluorescing side walls), it may be necessary to move the features manually to their final locations.

Tips:

- If the main grid cannot be moved far enough because it contacts the side of the image, move the subgrids instead. An easy way to select all the subgrids is to select the main grid, right click on the image, and select **Select All Subgrids On Grid** from the context menu. All the subgrids can then be moved by clicking and dragging them. The subgrids can be moved outside of the cells of the main grid without causing any problems.
- If the main grid template is to be used for multiple similar images, the template can be fine-tuned by measuring the X and Y offsets and entering the offsets in the main grid template. After the grid is in its final location, draw a selection rectangle from the upper left corner of the image to the upper left corner of the main grid and get the offsets from the status message at the bottom of the Odyssey window. Alternatively, the properties can be opened and another template created, thereby storing the current X- and Y-offsets and all other values in the new template.
- The **Save Grid As Template** command on the **Analyze** menu is also useful for saving grid templates in certain cases. The utility of this command is based on the fact that it saves everything about a grid

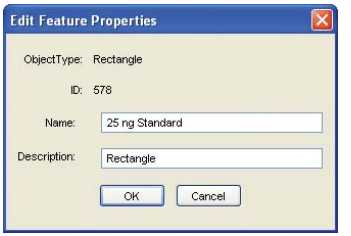
(i.e. much more information than is in the grid template). It saves everything about the grid, including any adjustments to individual features, grid rotation, etc.

Using the Auto Shape Tool

The purpose of the Auto Shape tool ( ) is to automatically create a feature that encloses fluorescence from irregularly shaped tumors or organs of small animals. The Auto Shape tool is disabled unless you have purchased a key to unlock the software for the Odyssey MousePOD™ Module.

Naming Features and Adding Annotations

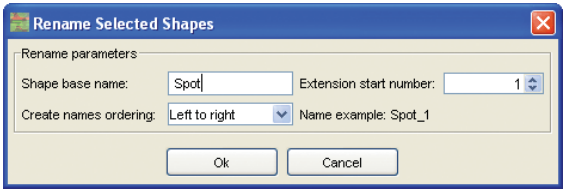
Each feature drawn on the image is automatically assigned an ID number by Odyssey software. To make a dot or band easier to identify on reports, a name and description can be assigned. To name a dot/band, select the feature that encloses it and click Properties ( ) on the toolbar. The name and description are retained with the feature.



Name or description annotations are not displayed unless they are enabled in the Application settings (see Chapter 11) in order for them to be displayed.

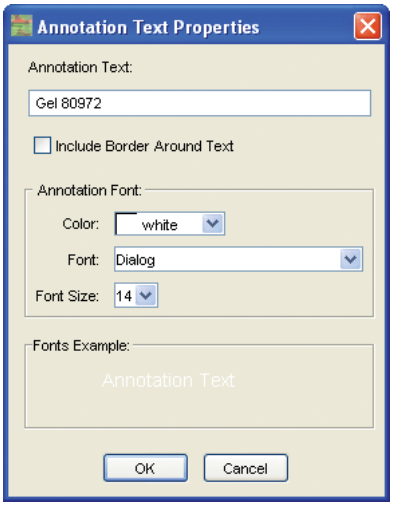
Renaming Multiple Features

Multiple features can be simultaneously renamed by selecting the features and choosing **Analyze > Rename Selected Shapes**. Features are named with a prefix (base name) and a numbered extension (see **Name Example** in the window below). Names are added to features in the order specified: left-to-right, right-to-left, top-to-bottom, or bottom-to-top.



Adding Text Annotations

To add an annotation, click the text tool () and then click once on the image where you want the upper left corner of the text to start.



Note: *If feature names or annotations are truncated at the edge of the image, use the Application Settings to extend the text area around the image as described at the end of Chapter 11.*

The Annotation Properties window can be used to enter a single line annotation, set the font properties of the text, or add a border to the text using the **Include Border Around Text** check box. The text properties can be changed at a later time by selecting the annotation and clicking the properties tool ( ).

Changing an Annotation

To change an annotation, click the annotation to select it and click the properties button ( ) to open the Annotation Text Properties window. Make changes and click **OK**.

Copying and Pasting an Annotation

To copy an annotation, click the annotation to select it, choose **Edit > Copy**, move the mouse cursor to the location where the annotation should be pasted, and choose **Edit > Paste**. A copy of the annotation will be pasted at the cursor position.

Rotating Annotations

After adding text annotations, annotations can be rotated using one of two methods. To change the rotation for all annotations on the image, choose **Settings > Application** and use the **Rotation** field in **Image View Features** to change the rotation to none, 45 degrees counterclockwise, or 90 degrees counterclockwise. Annotations can also be rotated by selecting them, right-clicking the image, and choosing the rotation from the popup menu. The popup menu allows each annotation to be rotated independently, rather than applying the same rotation to all annotations.

Other Annotations You May See

Either integrated intensity or concentration will be displayed after a feature is created, depending on the Application settings. If concentration values are displayed, the concentration is **n/a** (not assigned) until concentration standards are assigned and the concentration of unknown dots/bands can be calculated.

If the Application settings are set to display integrated intensity values, integrated intensity will be displayed immediately after the feature is drawn.

Hiding Annotations

If annotations obscure other image features, annotations can be temporarily hidden by clicking on the toolbar, or by choosing **View > Hide Annotations**. To view hidden annotations, click on the toolbar again, or choose **View > Show Annotations**.

Note: *To reduce screen clutter, use the Application settings to control which annotations are displayed for both selected and unselected features.*

Chapter 8: Quantification

Overview

Quantification can begin after a new analysis is started or an existing analysis is opened (Chapter 3). Before quantification, features must be drawn on the image (Chapter 7), or lanes and bands must be found (Chapter 5) for images with bands in lanes.

For scans with both 700 and 800 channel images, concentration standards in one channel cannot be used to quantify dots or bands in the other channel. Concentration standards for both dyes must be loaded and each image channel must be analyzed separately in Single Channel mode.

After all the features are drawn, concentration standards are identified by selecting the feature surrounding the standard and using the Concentration Standards window to enter the concentration. **Odyssey requires that concentration standards be added in order** (either lowest-to-highest or highest-to-lowest). Visually identifying all the concentration standards on the image before beginning will make it easier to add the standards in the correct order.

As soon as the concentrations of all standards are entered, the concentrations of all sample dots or bands are automatically calculated.

Quantification and Concentration Calculations

When a feature is drawn over fluorescence in an image, the image data within the feature are quantified immediately. This applies to features like rectangles or ovals that are explicitly drawn and to band markers that are automatically assigned when lanes are created.

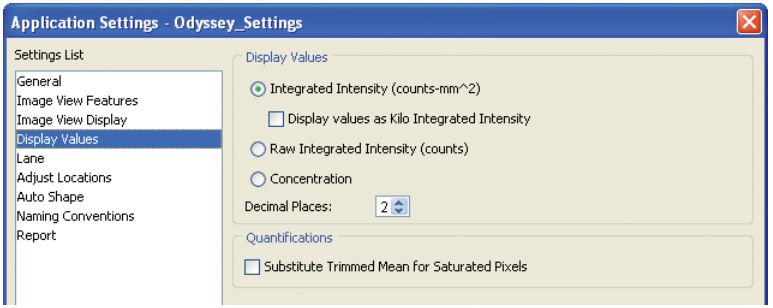
When a feature is quantified, **Integrated Intensity** is calculated. Integrated Intensity has also been referred to as **Pixel Volume** in other software. Integrated Intensity is the sum of the intensity values for all pixels enclosed by a feature, multiplied by the area of the feature (counts mm^2). Since background pixels should not be part of this calculation, background is calculated and subtracted. Several methods are available for calculating background. The calculation method is selected in the Application settings as described later in this chapter. Chapter 13 describes all of the calculations used in Odyssey software.

When the concentration of each standard is identified, a concentration value is assigned to the calculated integrated intensity for each standard. After the standards are all identified, concentration values of sample dots/bands are calculated from their integrated intensities using the interpolation method specified by the user.

Displaying Quantification Values

The Application settings determine how concentration and quantification values are displayed for band sizing or quantification on membrane scans.

To change how values are displayed, choose **Settings > Application** and select **Display Values** from the **Settings List**.



Quantification values can be displayed as integrated intensity (select **Integrated Intensity (counts-mm2**)), raw integrated intensity (select **Raw Integrated Intensity (counts)**), or concentration (select **Concentration**). The number of decimal places used when the concentration value is displayed can be changed using the **Decimal Places** field. Values are displayed both above a feature (if turned on) and in a tool tip as the cursor passes over a feature.

When **Display Values as Kilo Integrated Intensity** is selected, all integrated intensity values are divided by 1000 to make the values easier to read. This changes the display of integrated intensity values in all windows including the Grid Sheet window, ICW Analysis window, Details View, and any integrated intensity values displayed in the Image View window.

Note: *The Application settings control whether quantification values are displayed on the image (see Chapter 11).*

When **Substitute Trimmed Mean for Saturated Pixels** is selected, the calculated Trimmed Mean value (Chapter 133) is substituted for any saturated pixels enclosed in a given feature before quantification.

Raw integrated intensity values (in counts) should be used cautiously when comparing shapes, as detailed in Chapter 13. Raw integrated intensity values vary with resolution and the size of the feature drawn

on the image. Do not use raw integrated intensity values to compare features from images scanned at different resolutions. Similarly, image features drawn on the image must be exactly the same size if their raw integrated intensity values are to be compared.

Note: *For a band in a lane, the quantification value of the band can also be expressed as the band's percentage of the total integrated intensity of all bands in the lane (see Application settings at the end of Chapter 5).*

Entering the Concentration of Standards

Important: For scans with both 700 and 800 channel images, concentration standards in one channel cannot be used to quantify bands in the other channel. Concentration standards for both dyes must be loaded and each image must be analyzed separately in Single Channel mode.

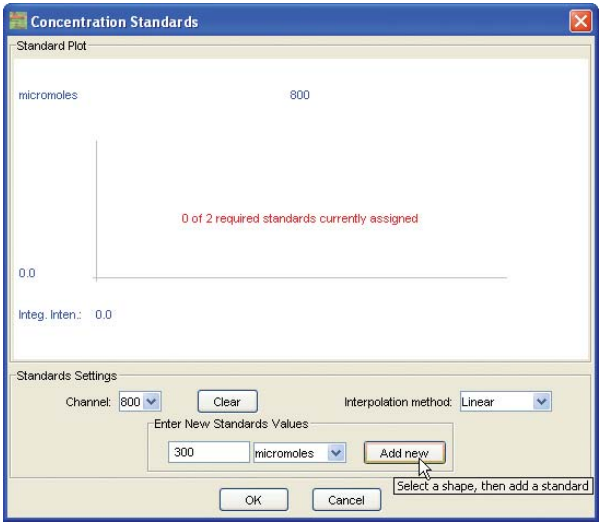
Concentration standards are identified using a shape tool (rectangle, circle, oval or freeform shape) to draw a feature that encloses the standard. Circles or rectangles in a grid can also be used as standards. On images where lanes have been added, the band markers surrounding the concentration standard bands can be selected and used for quantification.

Note: *Image background can be added as a "standard" by drawing a feature around an area of uniform background and adding it as a standard as described below.*

Important: *Concentration standards must be added in either ascending or descending order*. To help add the standards in the correct order, visually identify all concentration standards on the image before beginning.

After drawing a feature around a concentration standard, or selecting an existing feature like a band marker encompassing a concentration

standard, open the Concentration Standards window by choosing **Analyze > Concentration Standards** or click on the toolbar.



First, make sure the **Channel** (700 or 800) matches the image being analyzed. Next, enter the concentration for the selected standard in the **Enter New Standards Values** field, select the units, and click **Add New**.

Note: *Changing units changes the units for all standards on both images. The units need only be set once for each analysis.*

The first standard is now added. Continue adding the other standards by selecting a feature, entering the concentration in the Concentration Standards window, and clicking **Add New** (the Concentration Standards window can be left open). *The standards must be added in ascending or descending order.*

Setting the Interpolation Method

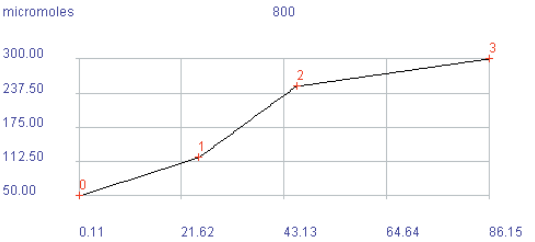
After the standards are entered, the interpolation method should be set and the standards plot reviewed for anomalous standards.

The concentration standards plot shows integrated intensity on the X-axis plotted against concentration on the Y-axis. For each dot or band of unknown concentration, Odyssey uses the integrated intensity of the dot/band and interpolates between the known values of the concentration standards to find the concentration of the unknown dot/band.

In general, the **Reciprocal Fit** interpolation method usually produces the best results. There are exceptions, however. If the plot of the standards is very linear, the **Linear** interpolation may give slightly better results. If the image has sample dots/bands that are a higher or lower concentration than any of the standards, use the **Log** interpolation method. Whenever there is a need to extrapolate beyond the known concentration standards, the **Log** interpolation method usually produces the best results.

Reviewing the Standards Plot

The standards plot should also be used to look for anomalous standards. Any standard that is out of position on the plot may need editing. For example, if a set of standards has a linear plot, but the straight line is broken by a standard that is too high or too low, the standard should be reviewed. *Standard #2* in the standards plot below appears to have a concentration value that is too high.



When reviewing the anomalous standards, make sure the standards are fully enclosed and that the feature drawn on the image is

centered over the fluorescence. Also make sure that the correct concentration has been assigned to the correct feature.

After examining the standards for one channel, use the Channel drop-down list to switch to the other channel and examine the plot of the concentration standards on the other image.

Changing and Deleting Concentration Standards

After a concentration standard is defined in the Concentration Standards window, it cannot be changed. To assign a different concentration value to a standard, all standards must be cleared and reassigned using the new values. Standards are cleared by clicking the **Clear** button in the Concentration Standards window.

Note: *Deleting a feature on the image that encloses a concentration standard does not delete the standard.*

Using the Details View for Background Verification

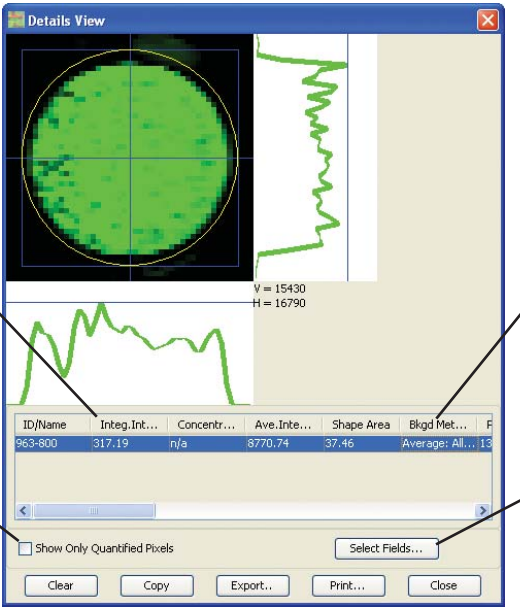
Having the Details View open when new features are drawn on the image gives immediate access to a variety of useful information about the newly quantified feature. Details View also provides an easy way to compare data values for various features on the image.

The Details View is opened by choosing **Report > Details View** or by clicking  on the toolbar. Details view can also be opened by selecting a feature, right-clicking the image, and choosing **Details View** from the pop-up menu.

With Details View open, an enlarged view of the band/spot is displayed each time a feature is added to the image. Details View is useful for verifying the location of the feature over the band/spot and the background calculation method. The feature shown surrounding

the band/spot should be examined to make sure it fully encloses the fluorescence on the image. On the image below, the circle feature fully encloses the spot and is roughly centered over it, as indicated by the blue crosshairs. If a feature needs to be moved, it must be moved on the image rather than in Details View. (Features can be nudged in one pixel increments using the arrow keys as long as the mouse cursor is over the selected feature.) To the right and below the image are curves that plot the intensity of the pixels below the crosshairs.

Any column in the Details View can be moved by dragging its header left or right to a new position.



*The background method used when the integrated intensity of this feature was calculated is listed in the **Bknd Method** column of the data table.*

*Click **Show Only Quantified Pixels** to turn all pixels black except for those that will be used for quantification.*

*Click **Select Fields** to change which fields are displayed in Details View.*

Each dot/band should be examined to make sure the background method was correct. In most cases one background method will be correct for most dots/bands on an image, but there may be a few that require recalculation using a different method.

As mentioned earlier, the sides of the blue rectangle in Details View are the horizontal and vertical boundaries of the feature on the image. The pixels on the outside perimeter of the blue rectangle are the pixels used for background calculations.

In the Details View shown above, the pixels just outside the blue rectangle on all four sides are empty background pixels. For this reason, either the **Average** or **Median-All** method (defined below) are appropriate. Both of these background calculation methods use the pixels surrounding all four sides of the blue rectangle.

Suppose, however, that there was another dot very close to the dot shown in Details View and that fluorescence from the second dot contacted the left side of the blue rectangle. In that case, the left side should not be used in the background calculation because the added fluorescence would increase the calculated background. The **Average** and **Median-All** methods would not be appropriate because they use all four sides. In this case, the **Median** method that uses only the top and bottom lines would be the appropriate background method. The **User Defined** background method could also be used.

Comparing Data Using Details View

Details View lists a variety of useful values, including concentration. With the Details View open, each time a feature is selected, the information for that feature is added to the table. To compare the concentration of two features, just select them and examine the values in the **Concentration** column. To clear the table, click the **Clear** button. A description of each column in Details View can be found in the field definitions for reports in Chapter 10. The calculation descriptions in Chapter 13 provide additional information.

The values shown in the table in Details View can easily be transferred to a spreadsheet for plotting or other operations. Click **Export** to create a text file (tab-separated values) containing a header that describes the scan and all the data in the Details View data table.

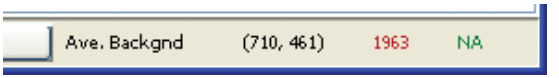
Click **Copy** to copy all the data in the Details View data table to the clipboard. Individual rows can be copied by Control-clicking the row(s) and pressing **Ctrl+c** to copy the data to the clipboard. To print the Details View table to the default printer, click the **Print** button.

Note: *Data can also be viewed, exported or printed using the menu choices on the* ***Report*** *menu (Chapter 10).*

Choosing the Background Calculation Method

Determining the Current Background Method

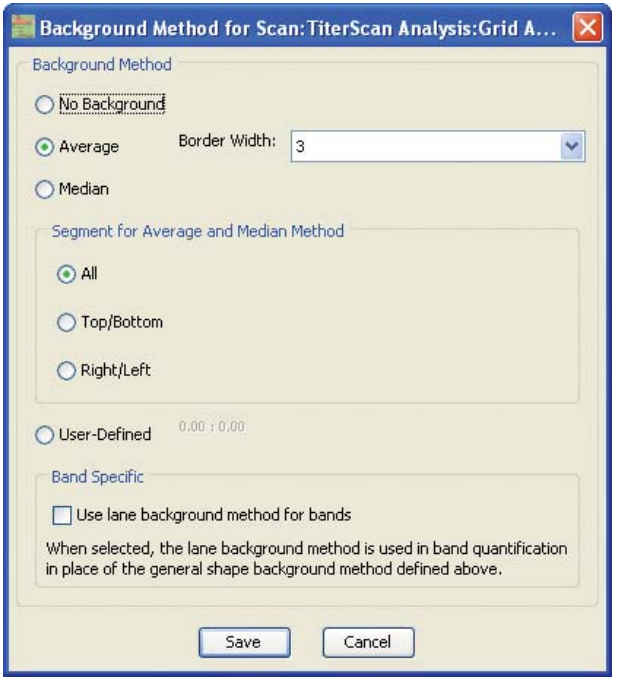
The background method is stored with each analysis and is listed in the tool tip that opens when the cursor is stopped over an analysis in the navigation pane. If an analysis is open, the current background method is always shown in the message line in the lower right corner of the main Odyssey window as shown below (Ave. Backgnd).



A message containing the *current* background method is displayed when an analysis is opened unless the message is turned off in the General Application Settings. Stopping the cursor over the background tool in the toolbar also displays the current background method in a tool tip.

Changing the Background Method

The background method sets the background calculation method for quantification. (See Chapter 9 for the background calculation method for In-Cell Western assays.) To change the background method for the current analysis, choose **Analyze > Background Method**.



Each of the background methods is briefly described below.

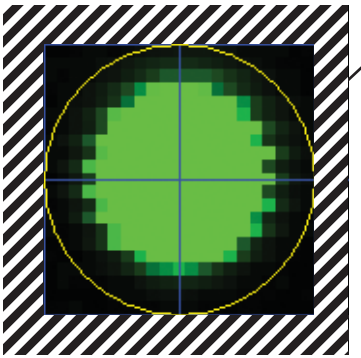
No Background

When **No Background** is selected, zero is used for the background. This is the best choice for applications such as In-cell Western assays that have their own background calculation method.

Average, Median, and User-Defined Background Methods

Both the **Average** and **Median** methods calculate the background using pixels around the perimeter of the area being quantified. The

Details View shows the feature drawn around the fluorescence and a blue bounding rectangle. The pixels used in the background calculation are the pixels between the blue rectangle in Details View and the edge of the image. The **Border Width** parameter in the Background Method window determines the width, in pixels, of the image outside the blue border that is used for background calculation.



To illustrate the location of the pixels used for Average or Median background methods, the pixels have been covered with diagonal lines.

When **Median** or **Average** is selected, background is calculated as the median value or average of the background pixels on the user-selected sides of the blue rectangle, respectively. After selecting **Median** or **Average**, choose which line segments of the perimeter rectangle to use in the calculation. If **All** is selected, pixels in all four sides of the blue rectangle are used. If **Top/Bottom** or **Right/Left** is chosen, only pixels in the two indicated sides of the rectangle are used.

User Defined: To use the **User Defined** background method, display both image channels and draw a feature on the image over an area of typical background. The feature will be added to both 700 and 800 channel images. With the channels still overlaid, select the feature, choose **Analyze > Background Method** and change the background method to **User Defined**. After clicking **Save**, any new features will

be quantified using the new user-defined background. The background feature is labeled 'background' to distinguish it from other features. To use more than one feature as background, select (**Ctrl + click**) all background features before opening the Background Methods window.

Note: *For some applications, such as in vivo imaging, background features are added to each image individually, rather than with channels overlaid.*

The background in the **User Defined** method is calculated as the average intensity of the pixels enclosed by the feature that is selected. To find out what the calculated background value is, open the Background settings again. The calculated background value is shown next to the **User Defined** radio button. If the scan is a two-channel scan, a value is shown for each channel (channel 700 followed by channel 800). The values are also shown in Details View.

Choosing Between Average, **Median**, or **User Defined**: The blue rectangle in Details View should be observed when determining which background method to use. If the pixels just outside the blue rectangle are dark background pixels, then either the **Average** method or the **Median** method with **All** boundaries will produce good results. If there are bands that are vertically close together with fluorescence in between them, it is best not to use the top and bottom segments in the background calculation. **Median** with **Right/Left** segments will produce good results. Similarly, when bands or dots are horizontally close together with fluorescence in between, the **Median** method with **Top/Bottom** will produce good results as long as the top and bottom line segments are over empty background. In cases where there is a lot of fluorescence from other bands/dots, neither the **Right/Left** or **Top/Bottom** lines may be usable. In this case, use the **User Defined** method as described above.

Using the Lane Background Method for Bands

The **Use Lane Background Method For Bands** check box is used when band markers within lanes are selected for quantification. When **Use Lane Background Method For Bands** is selected, the band background is calculated using the lane background profile (see Chapter 6 for information on lane profiles). The band background is calculated as the average intensity of the two points where the band boundary intersects the lane background profile (Chapter 5). The lane background profile can be displayed by selecting lanes and choosing **Lane > Lane Profile** to open the lane profiles window.

When **Use Lane Background Method For Bands** is selected and bands in lanes are being quantified, all other background settings (average, median, user-defined, or no background) are ignored. If there are both bands in lanes and other features being quantified, the lane background method will be used (if selected) for bands in lanes, and the other features will be quantified using average, median, user-defined, or no background, depending on which method is selected.

Note: *The background method used for each feature can be included in reports and is displayed in the data table in Details View.*

Requantifying After Changing Background Method

When changes are made to the background method, all existing features and band markers are automatically quantified again. Any new features or band markers added to the image will be quantified using the new background calculation method. A manual method to requantify features is also available. To recalculate existing features, choose **Analyze > Requantify**.

Quantification Using Grids

After all the features in the grid are properly positioned (Chapter 7), quantification can proceed as described earlier in this chapter. Concentrations are assigned to wells containing concentration standards by selecting the individual circle or square features and choosing **Analyze > Concentration Standards**.

Integrated intensities can be viewed immediately in the Grid Sheet (choose **Analyze > Grid Sheet**). After the concentration standards are assigned, concentration values are automatically calculated for all other features in the grid. Concentration values can also be viewed by generating a report (Chapter 10) that lists the concentrations for each feature.

Note: *Background calculations are set to No Background when a grid is applied to an image, so a background method should be selected before quantification.*

Chapter 9: In-Cell Western Module

Notice: Features described in this chapter are disabled unless you have purchased and imported a key to unlock the In-Cell Western Module.

Overview

It has been well documented that protein phosphorylation/dephosphorylation by kinases and phosphotases is a critical process regulating almost every aspect of life. Abnormal phosphorylation is a cause or consequence of major diseases like cancer, diabetes, and rheumatoid arthritis. An In-Cell Western assay has been developed to simultaneously detect both the phosphorylated protein and normalize for total protein (relative number of cells in each well). In-Cell Westerns are highly reproducible, sensitive, and linear over a wide dynamic range. Features distinguishing In-Cell Westerns for analysis of signal transduction from other currently available technologies include: (1) Near infrared probes with excitation/emissions at 700nm and 800nm minimize interference from auto-fluorescence of cells and plastic plates, and from chemical compounds and other materials, particularly potential drug candidates. (2) *In situ* detection to simplify sample handling and avoid the degradative effects of sample extraction. (3) Simultaneous assessment of two targets enables quantitative and accurate measurement of phosphorylation of one target because of data normalization with the another target. Furthermore, the Odyssey In-Cell Western software has been designed for the purposes of background subtraction, data normalization, and percentage response of each sample over positive controls.

Starting a New In-Cell Western Analysis

This section describes the operation of Odyssey software after samples have been prepared according to one of the ICW protocols in the *Odyssey Application Protocols*.

1) Place the microplate on the Odyssey scan surface and align it using the alignment guide. Only 96- or 384-well microplates can be analyzed with Odyssey In-Cell Western software module. (See Operator's Manual for tips on scanning microplates.)

2) Start the scan and select the **Microplate** preset or a similar preset.

3) When the scan is complete, click **Save** and accept the default scan and analysis names or enter new names.

Applying a Grid Automatically

Note: *Grids can also be placed manually as discussed in Chapter 7. Automatic grid alignment is only available in the In-Cell Western Module.*

For microplate scans, Odyssey software can automatically place grids on images as long as the images meet certain restrictions. To automatically place a grid, choose **In-Cell Western > Align Grid**. Select a grid template that matches the microplate and click **OK**.

Automatic grid placement is only available for images of 96- or 384-well microplates that have been scanned using Odyssey software 2.0 and above. Trying to automatically place grids on older images will generate at error message. Microplate images should be approximately 12 cm W x 8 cm H, and the microplate should be approximately centered in the scan area. For some images, such as those with weak fluorescence, grid alignment may fail, in which case the grid is placed according to the X and Y offset specified in the grid template.

On some images the grid may be placed correctly, but the circle or square features are too large or too small. This is possible because features are sized according to the size specified in the grid template. To get a better fit, delete the grid, change the size parameters in the grid template, and place a new grid. Images with a lot of background surrounding the microplate wells and images in which the microplate is not centered are also likely to cause problems with automatic grid finding.

Automatic Calculations

When a grid is applied, integrated intensity is calculated for each well (see Chapter 13). After integrated intensity is calculated, percent response is calculated for sample wells using the ICW template that was most recently used. Results can be viewed by choosing **In-cell Western > View ICW Analysis**. If the current microplate is setup differently than the most recently used ICW template (background wells, etc.), the ICW parameters can be changed and applied as described below.

Changing ICW Parameters for the Current Analysis

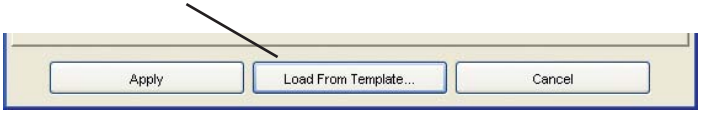
Choose **In-Cell Western > Change ICW Parameters** to change the parameters used in the ICW Calculations. The Change ICW Parameters window is used to switch to a different template, or just make temporary parameter changes that will be applied to the current analysis.

Applying a Different ICW Template

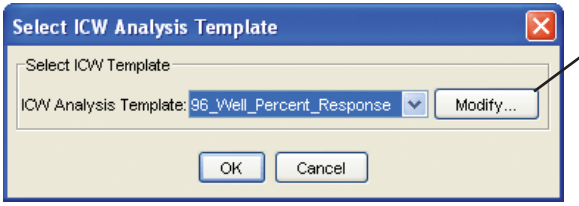
Use the following procedure to load a different ICW template and apply it to a grid.

1) Choose **In-cell Western > Change ICW Parameters**.

2) Click the **Load From Template** button at the bottom of the Change ICW Parameters window.



3) Select an ICW template from the list and click **OK**.



Modify the template as needed before applying it. (This is a permanent template modification.)

Note: *The new template must have he same number of wells as the grid on the image.*

4) Click **Apply** in the Change ICW Parameters window to apply the new template and recalculate the response data for each well using the well designations and calculation setup in the template.

Temporarily Changing the ICW Parameters

The Change ICW Parameters window is also used to change the parameters for the current ICW calculations without permanently changing a template. The Change ICW Parameters window is divided in to three tabbed panels that are used to specify how the microplate is loaded, which rows/columns to average, and which calculations to perform. Each of the tabbed panels is discussed below.

Well Types Tab

To change well designations, begin by selecting the type of well to mark (**Sample**, **100% Standard**, **Background**, or **Not Used**).

Move the cursor over a well on the well assignment grid and click to assign the selected well type to that well. Multiple well assignments can be made at once by clicking and dragging through a range of wells. When the mouse button is released, all wells within the selection rectangle change to the color assigned to the chosen well type.

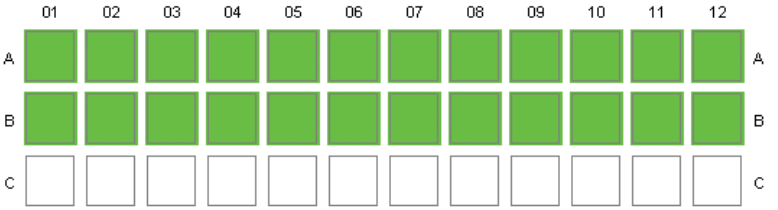
In some cases, it may be easier to mark sample wells last, since there are usually more sample wells than other types of wells. If **Sample** is selected, clicking **Assign Remaining** designates all wells that are **Not Used** as sample wells. To change all wells to **Not Used**, click **Clear All Rows**.

If more than one **Background** well is designated, all **Background** wells will be averaged for the ICW calculations (unless individual rows are being analyzed as described below). The same is true for **100% Standard** wells.

Well Links Tab

If two or more rows are exact duplicates of each other, the samples can be linked in the calculations to get an average response of all the duplicate samples. Suppose rows A and B are identically loaded and you want to average the response of similar samples in the two rows. After designating the well types, switch to the **Well Links** tab in the Change ICW Parameters window. Next, select all the wells in rows A and B by clicking the first well in row A, holding down the mouse button, and dragging through the last cell in row B. When the mouse

button is released, all the wells in both rows turn green to indicate they are selected.



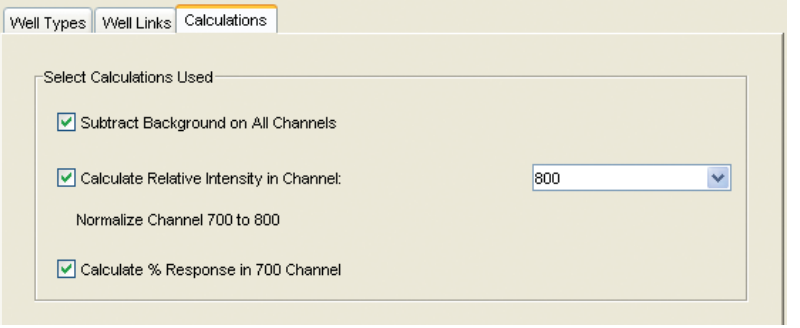
Select **Link Rows** and click **Create Link**. The green wells in the template turn black to indicate they are linked. Each individual link (A01-B01, A02-B02, etc.) is listed in the **Existing Links** list. During analysis, the linked wells in the **Existing Links** list will be averaged. The average will be used in ICW calculations rather than individual integrated intensity values. Only sample wells are averaged. (All **Background** and **100% Standard** wells are always averaged.)

Additional links between wells with identical samples can be added by continuing to select the wells and click **Create Link**. To link columns rather than rows, click **Link Columns** rather than **Link Rows**. To link a few wells that are not part of entire rows or columns, select the wells, select **Independent Link**, and click **Create Link**.

To dissolve a link, select the wells and click **Unselect** to change the well type from linked (green) to unlinked (white). Links can also be dissolved by clicking **Remove All** to remove all current row links, or selecting links in the link list and clicking **Remove Selected.** **Shift+click** and **Ctrl+click** are available for multiple selections.

Calculations Tab

After assigning any well links, click the **Calculations** tab. A complete description of the calculations can be found in Chapter 13.



When the ICW calculations start, background subtraction is performed on all wells for each channel if **Subtract Background on All Channels** is enabled. (Background subtraction is normally enabled.) During background subtraction, the integrated intensities of all wells designated as "background" in a given channel are averaged and subtracted from the integrated intensity of every well. References to integrated intensity throughout the rest of this discussion refer to the original integrated intensity minus background.

In a typical ICW analysis, the 700-channel might be used to detect phosphorylated proteins and the 800-channel used to detect total protein. Phosphorylated proteins in this case are probed with IRDye® 680-labeled secondary antibodies and total protein is detected using IRDye® 800CW-labeled secondary antibodies. The 800-channel, which is used to detect total protein, should be selected in the **Calculate Relative Intensity in Channel** field.

The next calculation compares integrated intensity values in the 800-channel (total protein) in order to find the well with maximum integrated intensity. All wells designated as **Sample** or **100% Standard** in the 800-channel are divided by the maximum integrated intensity to obtain the relative intensity of each well. The relative

intensity values will be between 0.0 and 1.0, which also indicates the relative number of cells in each well.

The relative intensity values from the 800-channel can now be used to normalize the integrated intensity values in the 700-channel, which is used to detect phosphorylated proteins. To normalize the 700-channel, the integrated intensity for each well in the 700-channel is divided by the relative intensity values from the 800-channel. This normalized value for each well is divided by the **100% Standard** of the 700-channel and multiplied by 100 to give a value that is the percentage response to the control in the **100% Standard**. If more than one well is designated **100% Standard** in the 700-channel, they are averaged before being used to calculate percentage response.

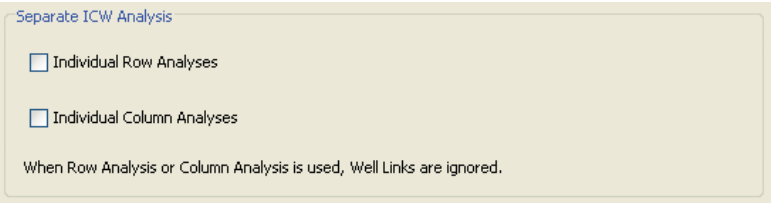
The calculation is slightly different if rows are linked in the well assignment window. When rows are linked, all the integrated intensity values for the linked wells in a given column are averaged. The average integrated intensity replaces the original integrated intensity values in all the linked wells in the column, resulting in them all having the same value.

Note: *If the* ***Calculate Relative Intensity in Channel*** *field is deselected for a two-channel scan, the percent response will be calculated for both channels with no normalization. For each channel, the relative intensity values are divided by the* ***100% Standard*** *and multiplied by 100.*

The **Calculate % Response** check box displays the channel(s) that will be used to calculate percent response. It cannot be deselected and has no other purpose but to provide information.

Analyzing Individual Rows or Columns

When the **Individual Row Analyses** and **Individual Column Analyses** check boxes are not checked, the ICW calculations are performed as described above.



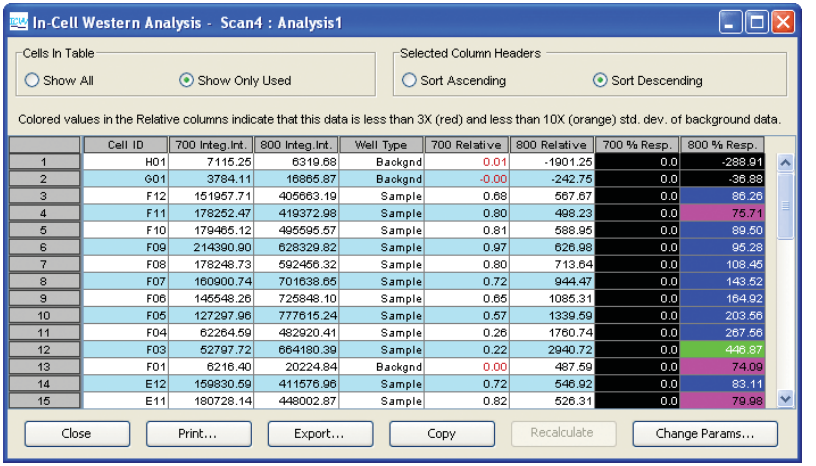
If either of these check boxes is checked, **%Response** is calculated using only the wells within a given row or column. For example, when **Individual Row Analyses** is checked, each row analyzed must have at least one **Background** well and **100% Standard** well. **%Response** for sample wells in the row is calculated using the **Background** and **100% Standard** well(s) in the same row as the sample wells. An error message will be displayed if **Background** and **100% Standard** wells have not been designated. If there is more than one **Background** or **100% Standard** well in a row, they are averaged before being used in the calculations, but they are not averaged with any **Background** or **100% Standard** wells outside the row. When individual rows are analyzed, wells between rows cannot be linked, so any links designated on the **Well Links** tab are ignored. When analyzing individual columns, the same concept applies: **Background** and **100% Standard** wells must be designated in each column to be analyzed and columns cannot be linked.

Applying the Changes

Click **Apply** in the Change ICW Parameters window to apply any parameter changes to the current analysis.

Examining the ICW Response Data

ICW response data can be viewed by choosing **In-Cell Western > View ICW Analysis**.



	Cell ID	700 Integ.Int.	800 Integ.Int.	Well Type	700 Relative	800 Relative	700 % Resp.	800 % Resp.
1	H01	7115.25	6319.68	Backgnd	0.01	-1901.25	0.0	-288.91
2	G01	3784.11	16865.87	Backgnd	-0.00	-242.75	0.0	-36.88
3	F12	151957.71	405663.19	Sample	0.68	567.67	0.0	86.26
4	F11	178252.47	419372.98	Sample	0.80	498.23	0.0	75.71
5	F10	179465.12	495595.57	Sample	0.81	588.95	0.0	89.50
6	F09	214390.90	628329.82	Sample	0.97	626.98	0.0	95.28
7	F08	178248.73	592456.32	Sample	0.80	713.64	0.0	108.45
8	F07	160900.74	701638.65	Sample	0.72	944.47	0.0	143.52
9	F06	145548.26	725848.10	Sample	0.65	1085.31	0.0	164.92
10	F05	127297.96	777615.24	Sample	0.57	1339.59	0.0	203.56
11	F04	62264.59	482920.41	Sample	0.26	1760.74	0.0	267.56
12	F03	52797.72	664180.39	Sample	0.22	2940.72	0.0	446.87
13	F01	6216.40	20224.84	Backgnd	0.00	487.59	0.0	74.09
14	E12	159830.59	411576.96	Sample	0.72	546.92	0.0	83.11
15	E11	180728.14	448002.87	Sample	0.82	526.31	0.0	79.98

If the data table is empty and a red status message is displayed indicating “Invalid ICW calculation parameters...”, the ICW calculation parameters (background wells, 100% response wells, etc.) have not been set up. To set up the ICW calculation parameters, click **Change Params** to open the Change ICW Parameters window (described earlier in this chapter).

Excluding Empty Wells

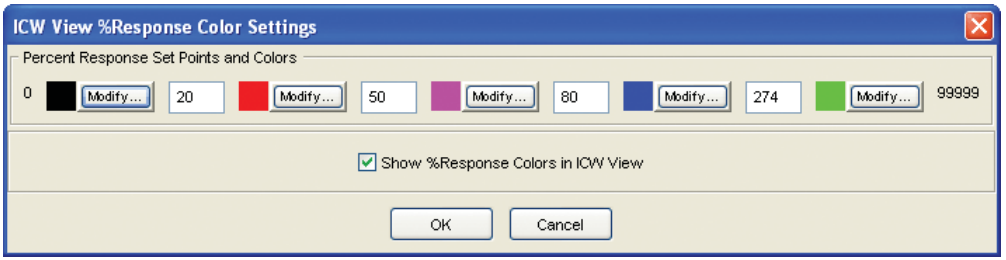
Initially, all wells are shown in the data table. Wells that were designated as "Not Used" in the well assignment window (listed as **Unused** in the **Well Type** column) can be excluded from the data table by selecting **Show Only Used**, rather than **Show All**.

Sorting Data

Start by selecting **Sort Ascending** or **Sort Descending** to set the sort order to ascending or descending, respectively. Next, click the column header of the data you want to use to sort the table.

Color-Coded Cells for Percent Response Values

The percent response values are color coded to visually distinguish cells with a given response. Table cells are initially colored according to the ICW View settings. To change response color codes, choose **Settings > ICW View**.



Colors are changed by clicking the corresponding **Modify** button. The response range associated with each color is changed by typing a new integer for the upper limit in the appropriate field. The lower limit of the range is always the upper limit of the previous range.

Deselecting **Show %Response Colors in ICW View** prevents percent response cells from being color coded.

Color-Coded Relative Intensity Values

If the relative intensity value for a given well is less than 3X the standard deviation of the background, the data will be listed in the ICW Analysis window with a font that is bold and red. Similarly, data values are orange when relative intensity is less than 10X standard

deviation of the background. These color-coded valued could be displayed in either the 700 Relative or 800 Relative column, depending on which channel is being used to calculate relative intensity. These color-coded values indicate the possibility of low quality data and the wells may need to be examined for errors (misalignment of the grid, etc.).

Recalculating Response Data

After viewing the response data, changes may need to be made and the data recalculated. The following changes require recalculation of the response data:

- Repositioning the grid on the image.
- Using the ICW parameters (**In-Cell Western** menu) to change well designations, linked lanes or how the calculations are performed (the channel used as the relative channel, etc.).

If changes are made that require the calculations to be performed again, the **Recalculate** button is activated.

Standard Deviation of Linked Wells

Standard deviations are calculated and reported for the integrated intensities of any wells that were linked during ICW setup. Values are reported in the **700 II Std Dev** and **800 II Std Dev** columns. For wells that are not linked, standard deviation values are listed as "n/a" (not applicable).

Exporting Response Data

Response data can be exported to a tab delimited text file by clicking **Export**. Use the standard file dialog to enter a file name and add any file name extension that your analysis program may require.

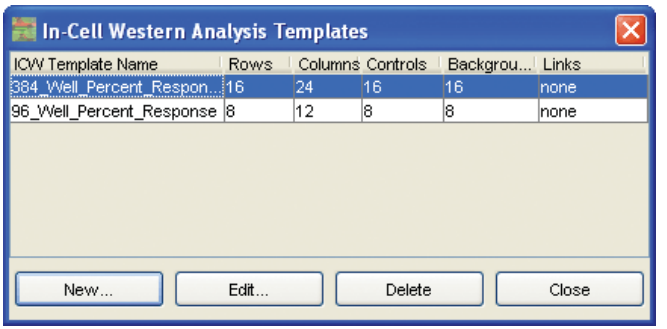
Data can also be copied or printed. Click **Print** to send the data to the default printer. To copy the entire data set to the clipboard, press **Ctrl+a** to select all cells, and then **Ctrl+c** to copy the data. Single columns are copied by clicking the column header to select the entire column and pressing **Ctrl+c** to copy the data. (Multiple columns can be selected by holding down the **Ctrl** key while clicking additional columns.)

Displaying Integrated Intensity in Kilo Units

The integrated intensity columns for the 700- and 800-channels may be displayed in standard units or divided by 1000 to make the numbers more readable. Chapter 8 describes how to use the Application settings (General) to change the display units.

Creating, Editing, and Deleting ICW Templates

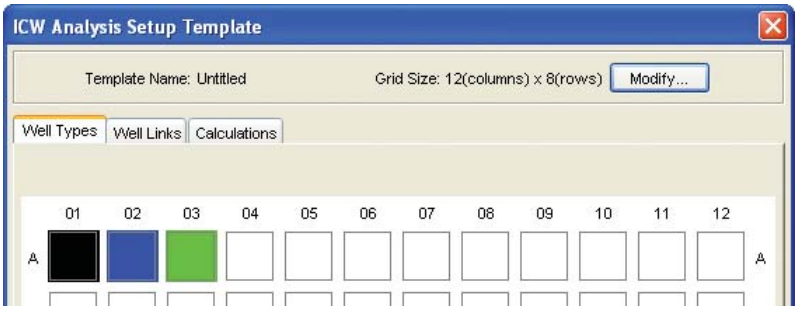
ICW templates can be created, edited or deleted by choosing **Settings > ICW Setup**.



The In-Cell Western Analysis Templates window lists all current templates and a few of their distinguishing parameters. To edit or delete a template, select the template in the list and click the **Edit** or **Delete** button, respectively. To copy a template, select the template,

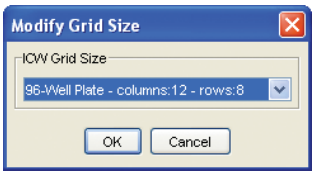
click **Edit**, change the template as needed, and click **Save As** to save the template under a different name.

To create a new template, click the **New** button.



The Setup Template window is nearly identical to the Change ICW Parameters window described earlier in this chapter except for some differences in the buttons at the top and bottom of the window.

The first thing to do in the Setup Template window is to check the grid size. Click the **Modify** button and choose the microplate that will be scanned: 96-, 384-, and 1536-well plates are supported, however 1536-well plates are not recommended for ICW assays.



The three tabbed panels in the center of the Setup Template window are identical to those describe earlier in *Changing ICW Parameters for the Current Analysis*. In the Setup Template window, the buttons at the bottom of the window are **Save** and **Save As**. One or both may be active depending on whether an existing template is being edited or a new template is being created. These buttons only save the

current template. The ICW parameters in the current analysis are unchanged.

Creating Reports for In-Cell Westerns

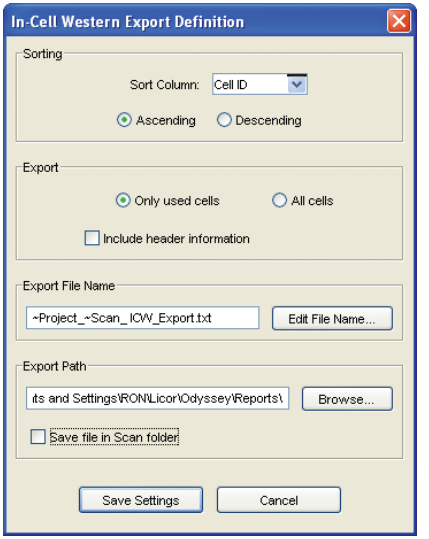
In-Cell Western reports are not available in the standard Odyssey Software. Standard reports are discussed in Chapter 10.

Printing and Saving Reports

To print an ICW report or save it to a file, choose **Print ICW** or **Export ICW** (respectively) from the **Report** menu. The form of the report is determined by the ICW report template, which can be changed as described below.

Changing the ICW Report Template

Choosing **Report > ICW Report Template** opens the ICW Export Definition window that can be used change the ICW report template.



The arrangement of data in a report is determined by choosing the column on which to sort and whether the data should be sorted in ascending or descending order. The **Export** section is used to output data for all cells in the table or to exclude cells that are marked as empty wells in the microplate. When **Include Header Information** is selected, information about the project, scan, and analysis is included in the beginning of the file. If **Include Header Information** is selected, the report will also include Z'-factor values if Z'-factor calculations are enabled.

The name for the export file can be entered using one of the two methods described below. This file name will be the default file name listed in the Save File window when the report is exported.

- Enter a simple file name in the **Export File Name** field.

- Click the **Edit File Name** button to create a file name that includes text, such as the analysis name, that is added automatically when the file is created. Names that are generated automatically save time when scanning multiple plates.

  In the file name above, the text "ICW Data From" is followed by two auto-entered text blocks (analysis name and date). Auto-entered text blocks can be separated by entering underscore characters, spaces, or other text as needed. An example of how a file name might look is shown in the **Example** field.

  Text that can be automatically inserted in the file name includes the **Date**, **Time**, **Project** name, **Scan** name, **Analysis** name, and organization name (choose **Settings > Application** and select **General** to enter an organization name). To insert a placeholder for the auto-entered text, select the type of text to enter from the drop-down list and click **Insert**. Place holders always start with a tilde character.

  Click **OK** when finished entering a title.

The directory path sets the default directory where the report file will be saved. To always store the report in the same directory as the scan, click **Save File in Scan Folder** and the path will be set automatically. A path can also be entered by typing it in the **Export Path** field or by clicking **Browse** and selecting a path from a file selection window. The path entered will be the default directory shown in the Save File window that is opened when exporting a report. Make sure to enter

the **Export File Name** and **Path** in such as way as to create a unique file for each scan, unless there is a reason to overwrite the export file with every new scan.

When finished making changes to the ICW report template, click **Save Settings** to save the report template.

ICW Export Settings

Another way to display the ICW Export Definitions window described in the previous section is to choose **Settings > ICW Export**.

Assay Optimization With Z′-factor Calculations

Calculating the Z'-factor for an assay produces a statistically derived, dimensionless value that can be used to characterize the quality of an assay during assay optimization and validation. The Z′-factor indicates whether the assay has sufficient dynamic range and low enough data variability to generate meaningful data. See the calculation descriptions in Chapter 13 to learn how Z'-factor is calculated.

Z'-factor values are always 1 or less as described below.

- $Z' = 1$ is an ideal assay.
- $1 > Z' \geq 0.5$ indicates a high quality assay exhibiting a wide separation between signals for positive and negative controls, and low data variability.
- $0.5 > Z' > 0$ may indicate a low quality assay with marginal distinction between signals for positive and negative controls, and higher data variability. However, an acceptable Z'-factor target value should be determined prior to performing final validation of an assay. An assay with a relatively low number of data points, such as the number obtained from a 96-well plate, may produce

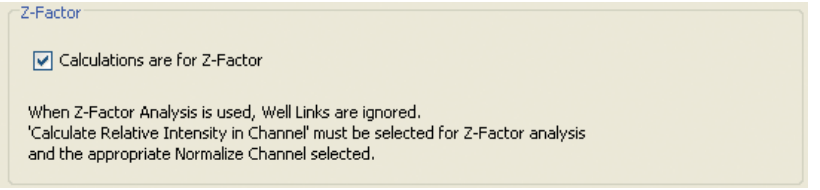
a Z'-factor value less than 0.5 but still be considered a good quality assay if the same value is acheived between different plates run on different days.

- Z' ≤ 0 indicates unreliable data.

A Z'-factor value that is close to zero or negative indicates assay conditions have not been optimized or the assay is not capable of generating meaningful data.

Enabling Z'-Factor Calculations

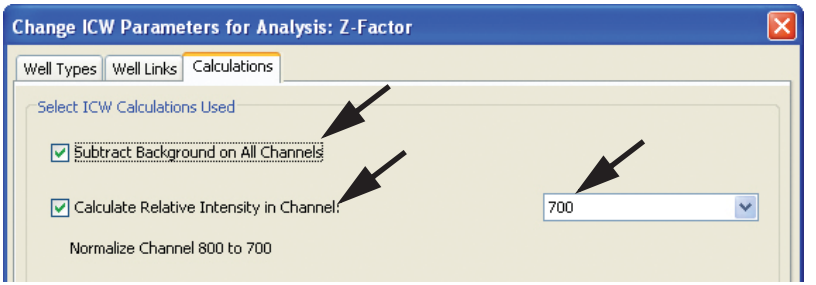
Z'-factor calculations are enabled by choosing **In-Cell Western > Change ICW Parameters** and switching to the **Calculations** tab. At the bottom of the **Calculations** tab, select **Calculations are for Z-Factor**.



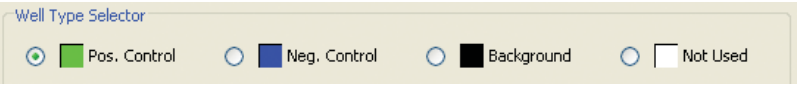
When **Calculations are for Z-Factor** is selected, In-Cell Western calculations are not performed and any well links are ignored if links are defined on the **Links** tab of the Change ICW Parameters window.

If background wells are loaded, make sure **Subtract Background on All Channels** is selected. When using Z'-factor calculations, one image channel should always be used for normalization in order to compensate for variations in cell number between wells. Make sure

Calculate Relative Intensity in Channel is selected and the correct channel for normalization is chosen.



Next, switch to the **Well Types** tab in the Change ICW Parameters window. For Z'-factor calculations, each well should be designated as a positive control, negative control, background well, or not used. To change well designations, begin by selecting the type of well to mark from the **Well Type Selector**.

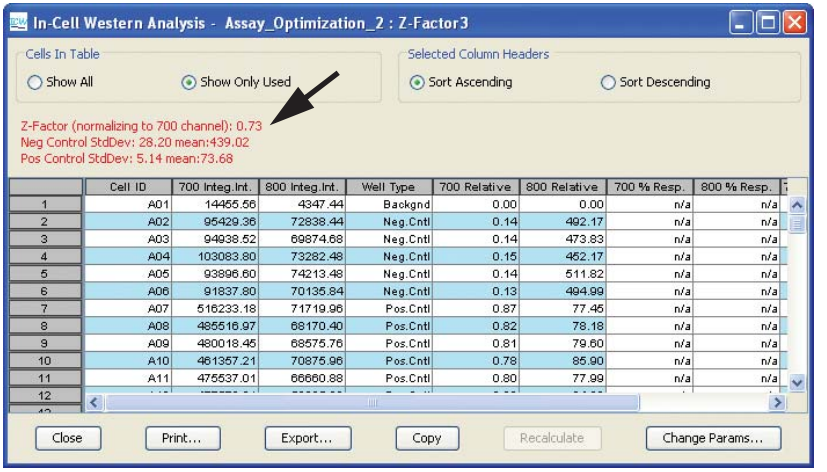


Move the cursor over a well on the well assignment grid and click to assign the selected well type to that well. Multiple well assignments can be made at once by clicking and dragging through a range of wells. When the mouse button is released, all wells within the selection rectangle change to the color assigned to the chosen well type. If **Assign Remaining** is clicked, any wells that are **Not Used** will be assigned the currently selected well type. Clicking **Clear All Rows** changes all wells to unused.

Viewing Z'-Factor Values

Choose **In-Cell Western > View ICW Analysis** to see the calculated Z'-factor value. The Z'-factor is listed in red above the data table in the View ICW Analysis window. Standard deviation and mean of the positive and negative controls are also listed. Percent response fields

in the data table contain "n/a" since ICW values are not calculated during Z'-factor calculations.



	Cell ID	700 Integ.Int.	800 Integ.Int.	Well Type	700 Relative	800 Relative	700 % Resp.	800 % Resp.
1	A01	14455.56	4347.44	Backgnd	0.00	0.00	n/a	n/a
2	A02	95429.36	72838.44	Neg.Cntl	0.14	492.17	n/a	n/a
3	A03	94938.52	69874.68	Neg.Cntl	0.14	473.83	n/a	n/a
4	A04	103083.80	73282.48	Neg.Cntl	0.15	452.17	n/a	n/a
5	A05	93896.60	74213.48	Neg.Cntl	0.14	511.82	n/a	n/a
6	A06	91837.80	70135.84	Neg.Cntl	0.13	494.99	n/a	n/a
7	A07	516233.18	71719.96	Pos.Cntl	0.87	77.45	n/a	n/a
8	A08	485516.97	68170.40	Pos.Cntl	0.82	78.18	n/a	n/a
9	A09	480018.45	68575.76	Pos.Cntl	0.81	79.60	n/a	n/a
10	A10	461357.21	70875.96	Pos.Cntl	0.78	85.90	n/a	n/a
11	A11	475537.01	66660.88	Pos.Cntl	0.80	77.99	n/a	n/a
12								

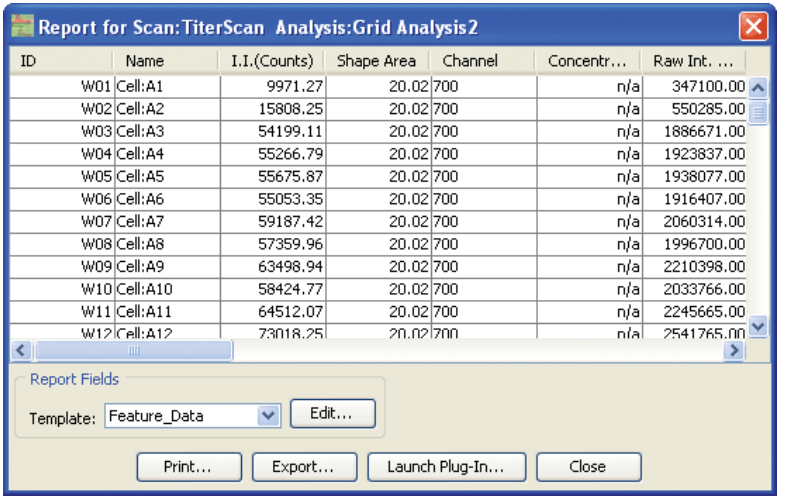
If the ICW export settings (**Settings > ICW Export**) are configured to include header information in reports, the Z'-factor values will be included in reports generated by clicking **Print** or **Export**.

Chapter 10: Reports and Data Export

This chapter discusses the standard reporting features of Odyssey software. ICW reports, which are available in the optional In-Cell Western Module, are discussed in Chapter 9.

Report Table View

Analysis data for *selected* features (grids, etc.) can be viewed or exported in table format by choosing **Report > Report View**.



The **Report View** menu choice is disabled until image features are selected and only data for selected features are displayed in the table. The report template, selected in the **Template** drop-down list, deter-

mines the fields displayed in the table, the sort order, and the default export path. If the current template is not what is needed, change the template by clicking **Edit** (which is exactly the same as choosing **Report > Report Field Templates**).

Click **Print** to print the data to a printer of your choice, or click **Export** to save data in a tab delimited text file. See *Printing and Exporting Reports* below for details.

The **Launch Plug-in** button displays a list of available plug-in reports. After selecting a plug-in and clicking **Launch**, data are transferred from the Report View to the plug-in and the plug-in performs the task it was designed for.

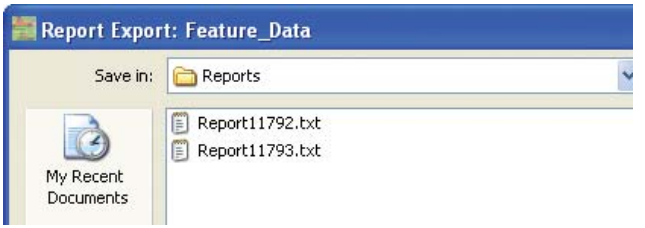
Consult *Plug-in Reports* below for troubleshooting tips regarding plug-ins.

Printing and Exporting Reports

The **Report > Print** and **Report > Export** menu choices are functionally the same as opening the Report View and clicking the **Print** or **Export** button, respectively.

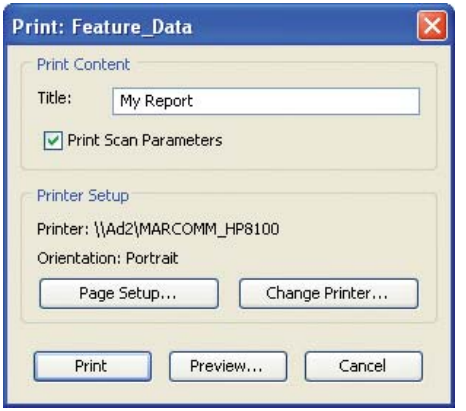
Important: *Both the **Print** and **Export** menu choices on the **Report** menu format data according to the current report template, which is selected in the Report View.*

When **Report > Export** or **Report > Print** is chosen, the name of the current report template is shown in the title bar of the window that is opened (Feature_Data is the current template shown below).



Printing Reports

Choose **Report > Print** to open the print window.



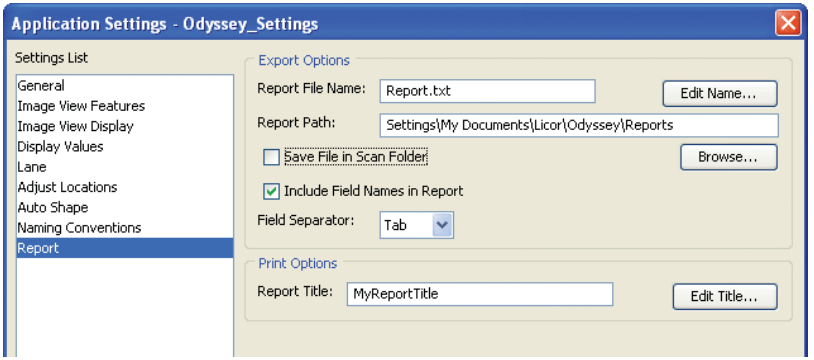
The default report title in the **Title** field is controlled by the Report Application Settings, but the title can be changed as needed. The scan parameters (resolution, etc.) can be appended to a report by selecting **Print Scan Parameters**. **Page Setup** (page orientation) and **Change Printer** are standard printer functions, and **Preview** shows a facsimile of the page that will be printed.

Exporting Report Files

When **Report > Export** is chosen, a standard save file window is opened and the default path is the **Report File Name** and **Path** in the Application Settings (**Settings > Application > Report**). The field separation character (tab, etc.) used in the file is also determined by the Report Application Settings.

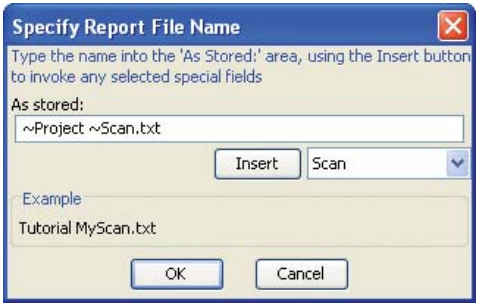
Report Settings

The report settings are opened by choosing **Settings > Application** and selecting **Report** from the **Settings List**.



The **Export Options** are all related to creation of data files. The **Report File Name** can be entered directly in the **Report File Name** field or a name can be created by clicking **Edit Name**. Names created

using **Edit Names** can contain placeholders for text such as project name, scan name, etc. The file name is assembled automatically when the report is exported.



In the title above, two auto-entered text blocks (project name and scan name) are followed by a file name extension (.txt). Auto-entered text blocks can be separated by commas, spaces, or other text as long as the resulting file name remains a legal file name. An example of how the file name might look is shown in the **Example** field.

Text that can be entered automatically includes the date, time, analysis name, project name, scan name, organization name, and report name. (Choose **Settings > Application** and then **General** to enter the organization name.) To insert a text placeholder at the cursor position, select the type of text to enter from the dropdown list and click **Insert**. Placeholders always start with a tilde character. Click **OK** when finished entering a title.

The **Report Path** (location of the exported file) can be entered directly in the **Report Path** field, or by clicking **Browse** to browse for the path using a standard file selection window. All exported reports will be stored in the same folder using the selected path. To store exported report files in the same folder as the scan, rather than a central location, select **Save File in Scan Folder**.

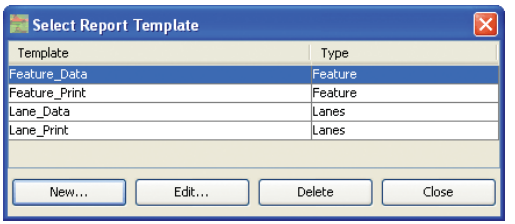
Analysis programs have varying requirements for input data files. **Include Field Names in Report** can be deselected if the analysis program does not allow field names at the top of each column. A comma, tab, or space can be chosen as the **Field Separator** character. The separator character is used by analysis programs to separate data values into columns. The separator character is also important for plug-ins since this character is automatically used as the field separator. Any plug-in designed for Microsoft Excel, for example, should use a tab as a field separator.

The **Report Title** can be entered as text, or a combination of user-entered text and text that is automatically entered by Odyssey software when the report is created. To use auto-entered text blocks, click **Edit Title** and use the **Insert** button to add placeholders for auto-entered text as described above for the report file name.

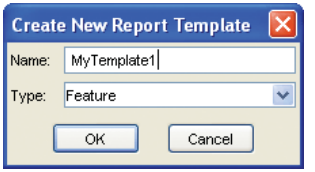
After configuring the report settings, click **Save** to store the new report settings. Note that these settings are saved only for the current Application settings file.

Creating Report Templates

Report templates are used for printed reports and for creating the data table in the report view. The current template can be selected or edited in the report view. Report templates can also be created, edited or deleted by choosing **Reports > Report Field Templates**.

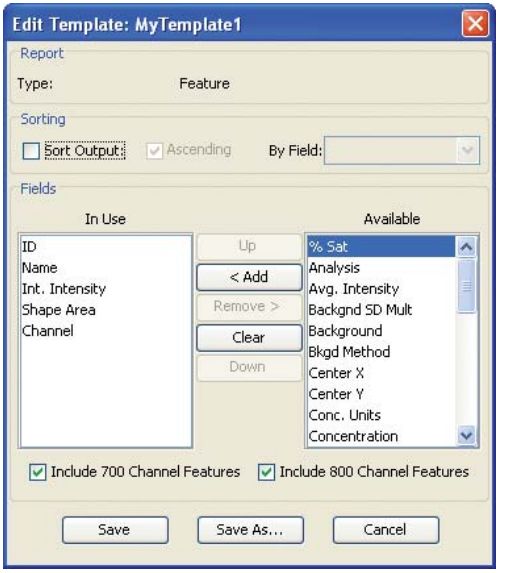


Click **New** in the Select Report Template window to start a new template.



Enter a name for the template in the **Name** field and select the template type. Select **Feature** from the **Type** drop-down list to create a template for selected features or **Lane** if the new template is for a report on lanes. Click **OK** to continue.

Both Feature reports and Lane reports are created the same way. The only difference is the fields that can be included in the report. The report type is always listed in the **Type** field at the top of the Edit Template window.



Choosing Fields to Include in the Report

Use the **Fields** area at the bottom of the Edit Template window to select the fields to include in the report. Definitions for each of the fields are given later in this chapter.

The **In Use** list shows fields that will be included in the report. By default, several fields are automatically added to the **In Use** list when the Edit Template window is opened. These fields can all be removed and added back to the **Available** list by clicking the **Clear** button. Fields in the **In Use** list can also be removed one at a time by selecting the field in the list and clicking **Remove**. To add a field, select the field in the **Available** list and click **Add**. To change the order in the **In Use** list and the report, select a field and click **Up** or **Down** to move the selected field.

Use the **Sort Output** check box to choose whether to sort the report data. If **Sort Output** is selected, you can choose the field by which records will be sorted and whether the order is ascending or descending. If the **Ascending** check box is selected, the order will be ascending, otherwise the sort order is descending. Use the **By Field** drop-down list to choose the field on which to sort.

By default, data for both image channels are included in reports. Deselect either the **Include 700 Channel Features** or **Include 800 Channel Features** check box to include only one image channel in the report.

Saving the Template

After editing the template, click **Save** to save the finished template.

Field Definitions

Feature Reports

Field	Definition
ID	ID numbers are automatically assigned to every feature (lane, circle, square, etc.)
Name	User-supplied name entered in the properties.
Description	User-supplied description entered in the properties.
Lane Name	User-entered name for the lane (entered in the Properties).
Channel	Name of the channel (700 or 800) in which the feature is found.
Concentration	Calculated concentration of the feature.
Conc. Units	Units selected in the Concentration Standards window.
Raw Int. Intensity	See description in Chapter 13.
Int. Intensity	See description in Chapter 13.
Avg. Intensity	Average intensity of all pixels that comprise the feature. See description in Chapter 13.
Peak Intensity	Highest intensity value of all pixels that comprise the feature.
Minimum Intensity	Lowest intensity value of all pixels that comprise the feature.
Pixels	Number of pixels in the feature.
Shape Area	Area of a feature in mm^2. Area is the number of pixels multiplied by the area per pixel. Area per pixel, $a=(\text{resolution} \times 10^{-3})^2$.

Field	Definition
Width	Width of an imaginary bounding rectangle around the feature.
Height	Height of an imaginary bounding rectangle around the feature.
Shape	Shape of a feature (circle, rectangle, etc.).
Ref Point X	X-coordinate value of the upper left corner of the feature.
Ref Point Y	Y-coordinate value of the upper left corner of the feature.
Center X	X-coordinate value of the center of the feature.
Center Y	Y-coordinate value of the center of the feature.
Background	Background value calculated for the feature. See description in Chapter 13.
Bkgd Method	Background calculation method used to calculate background. See Chapter 8.
Project	Name of project containing the feature.
Scan	Name of scan containing the feature.
Analysis	Name of analysis containing the feature.
MW	Molecular weight of a band.
MW w/units	Same as MW with units label.
%Sat	Number of saturated pixels divided by the total number of pixels, multiplied by 100 to give a percentage.
Probability	See description in Chapter 13.
Trimmed Mean	See description in Chapter 13.
SN Ratio	Signal-to-Noise Ratio. See Chapter 13.

Field	Definition
Std. Dev.	Standard Deviation of background pixels.
Background SD Multiplier	See Odyssey In vivo Imaging Guide Chapter 3. Values in this field are irrelevant unless the feature was created with the Auto Shape tool.
Normalized I. I.	Normalized integrated intensity. See Chapter 13.

Lane Reports

Field	Definition
ID	ID numbers are automatically assigned to every feature (lane, band, etc.)
Name	User-entered name for a band (entered in Properties).
Number	Band number automatically assigned by Odyssey.
Lane Name	Name for the lane (entered in the Properties).
Description	Description entered in the object properties by the user.
# Bands	Total number of bands in the lane.
Channel	Name of channel (700 or 800) where lane is found.
Concentration	Calculated concentration of a band.
Conc. units	Units selected in the Conc. Standards window.
MW	Molecular weight of a band.
MW w/units	Same as MW with units label.
Background	Background value calculated for the feature. See description in Chapter 13.
Bkgd Method	Background calculation method used to calculate background. See Chapter 8.
Rf	Relative mobility (Rf) is a measure of the distance a band has migrated as a percentage of the total lane length. Rf values are therefore a decimal value between 0 and 1.
Pixels	Number of pixels in the feature.
Shape Area	Area of a band in mm^2. Area is the number of pixels multiplied by the area per pixel. Area per pixel, $a=(\text{resolution} \times 10^{-3})^2$.

Field	Definition
% Sat	Number of saturated pixels divided by total number of pixels, multiplied by 100 to give a percentage.
Project	Name of project containing the lane.
Scan	Name of scan containing the lane.
Analysis	Name of analysis containing the lane.
% Total I.I.	Percent of total integrated intensity of all bands in a lane.
Raw Int. Intensity	See description in Chapter 13.
Integrated Intensity	Integrated intensity.
Normalization Factor	A multiplier between 0 and 1 used to normalize each lane in the reference channel.
Normalized I.I.	Normalized Integrated Intensity.

Plug-in Reports

A variety of plug-in reports can be accessed via the **Plug-in** menu. A description of each plug-in is given below.

Plug-in Name	Plug-in Type	Plug-in Description
Feature Data Report	Report	Sends data for all selected features to Excel. The data sent for each feature is dictated by a report template named Feature_Data, which can be modified to change the report. Only data for selected features are included in the report.
Microsoft Excel for Grids	Grid	Sends grid data to Excel. Choose **Report > Grid Fields Plug-in Template** to control which data are exported.
Microsoft Excel for ICW	ICW	Sends response data and integrated intensities to Excel using the current ICW report template.
Microsoft Excel for features	Shape	Exports all data for all features to Excel.
ReagentsWebLink	Command	Executes a command line statement that launches Internet Explorer and sets the page to the Odyssey reagents page at www.licor.com.

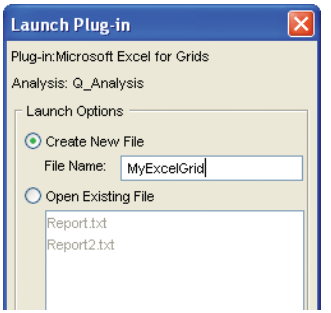
Launching Plug-in Reports

Plug-ins are launched by selecting them from the **Plug-in** menu. The example below shows the procedure for using the **Microsoft Excel for Grids** plug-in. Any user-created plug-in reports will follow the same general procedure.

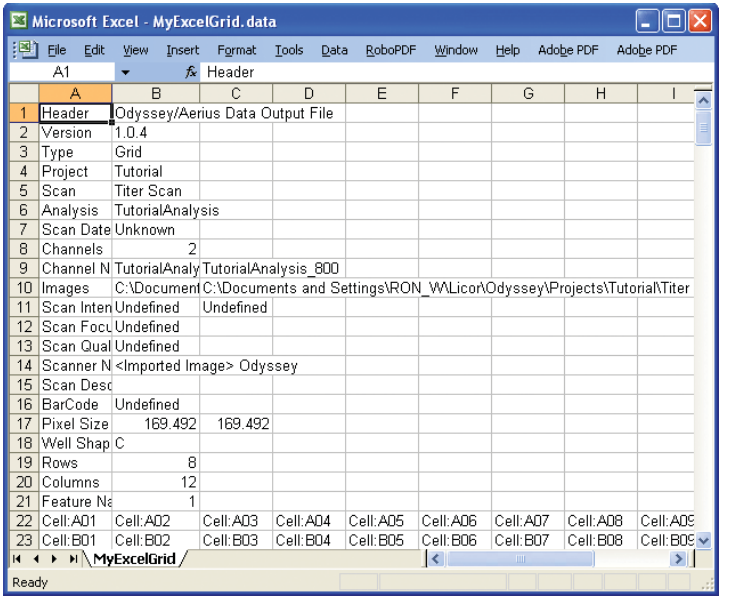
1) Open the analysis that has the grid data to export.

2) Choose **Plug-in > Microsoft Excel for Grids**.

3) In the Launch Plug-In window, select **Create New File** and enter a file name. The current analysis name from the Odyssey scan is entered as the default file name. An existing file can be overwritten by selecting **Open Existing File** and selecting a file from the file list.



4) Click **OK**. The Microsoft Excel program should start and the grid data should be displayed in a new worksheet.

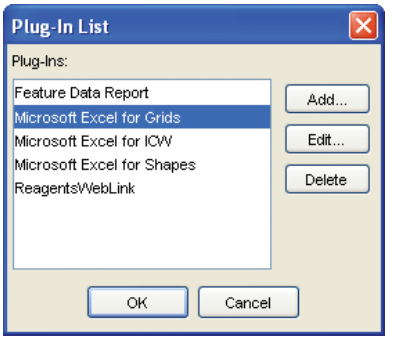


See *Troubleshooting Plug-in Reports* below if the program does not execute properly.

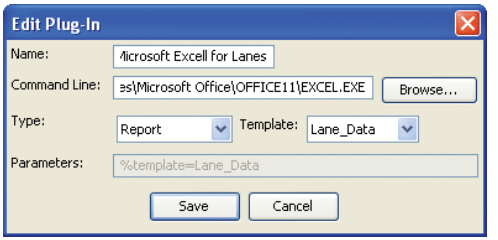
Editing Plug-in Reports

Though most parameters for plug-in reports will not need to be edited, plug-in reports such as those for Microsoft Excel include a path to the application (*.exe file) that may require editing for new versions of Microsoft Office.

To edit a plug-in report, choose **Plug-in > Plug-in Settings**.



In the **Plug-in** list, select the plug-in report that needs to be edited and click **Edit**. The **Name** field is the only field that does not directly influence the operation of the plug-in.



The **Command Line** should specify the program to start when the plug-in is launched. The program and path can be specified by clicking **Browse** and using the file selection window to find the program. Any program that accepts a data file parameter when launched from a command line can be used in the plug-in. The program must be able to accept data separated by tabs, spaces, etc.

The **Type** list is used to specify what the plug-in does and how it interacts with the **Command Line** and **Parameters**. Each plug-in type is described below:

- **Grid** exports a matrix of values for each field specified in the Grid Fields Plug-in Template (edit by choosing **Report > Grid Fields in Plug-in Template**). Data are arranged in a matrix that matches the features in the grid. For example, if integrated intensity is one of the exported fields for a grid that matches a 96-well microplate, integrated intensity values for each feature in the grid are arranged in an 8 x 12 matrix, followed by another 8 x 12 matrix for some other specified field. Data are only exported for fields that are selected in the Grid Fields Plug-in Type Template window.
- **Report** is used for plug-in reports. When **Report** is selected, the **Template** list is activated so a report template can be selected. Only data for the fields specified in the report template are exported. When a report template is selected, the **Parameters** are automatically modified to use the new template.
- **Shape** exports a complete data set for each selected feature (shape). (The complete data set is all fields available in reports.)
- **ICW** exports the complete In-Cell Western data set that is displayed in table form when **In-Cell Western > View ICW Analysis** is chosen. **Parameters** should always be set to %data for ICW exports.
- **Command Only** is the same as typing the contents of the **Command** field and the Windows command prompt, followed by the parameters in the **Parameters** field. To understand this plug-in type, examine the default ReagentsWebLink plug-in. The **Command** field contains the path to Internet Explorer and the **Parameters** field is the URL for the page that Internet Explorer is supposed to open when it launches.

The **Parameters** field should normally be one of the following:

- %data for plug-ins that export data.
- %template= for report plug-ins, where the text that follows the equal sign is the name of a valid report template.
- Parameters that must follow the command for a **Command Only** plug-in.

Important: Select the features or grid for which to export data before running the plug-in. Only data for selected features are exported to plug-ins.

Adding and Deleting Plug-ins

Plug-ins are created by choosing **Plug-in > Manage Plug-ins** and clicking the **Add** button in the Plug-in List window. The **Name**, **Command Line**, **Parameters**, **Type**, and **Template** can be entered as described above.

When plug-ins are no longer needed, they can be deleted from the list in the Plug-in List window by selecting them and clicking **Delete**.

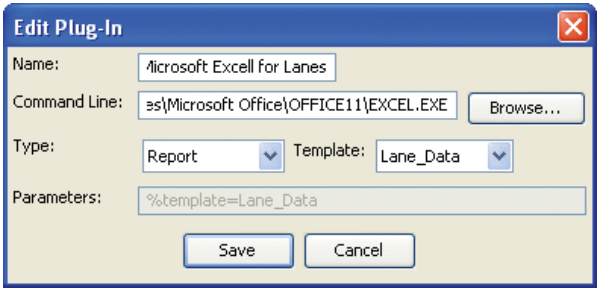
Troubleshooting Plug-in Reports

The most common problem is that the path to the analysis program receiving the data (Excel, etc.) is not correct. If the program is not found when the plug-in is launched, an error message is displayed. To change the path, use the following procedure:

1) Choose **Plug-in > Manage Plug-ins**.

2) Select the plug-in to be edited from the list and click **Edit**.

3) Make sure the path in the **Command Line** field is the correct path. If not, use the **Browse** button to find the program. (Programs generally have a .EXE file name extension (EXCEL.EXE, etc.).



4) Click **Save** and run the plug-in again.

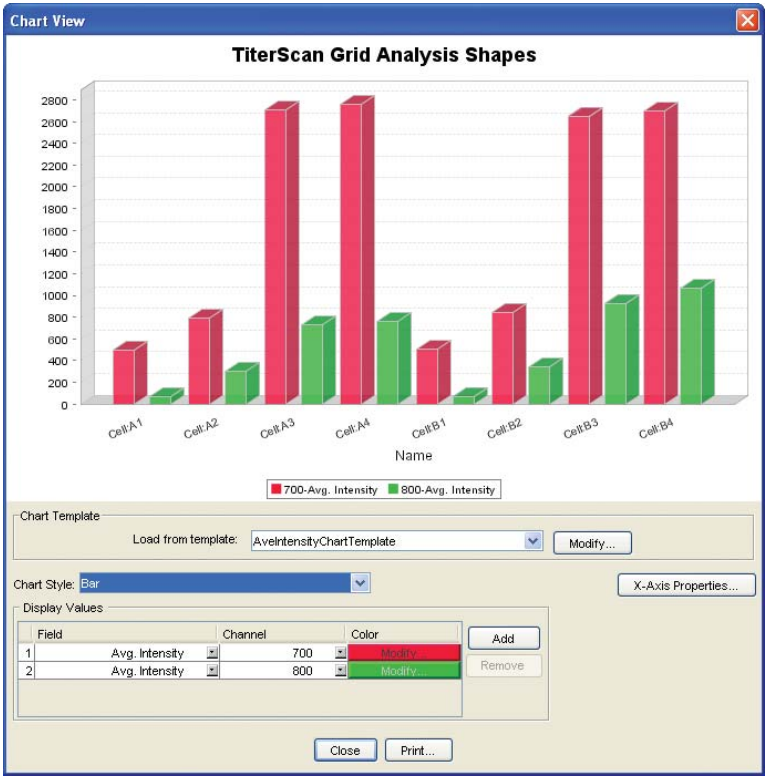
Another common problem is having the wrong **Field Separator** character selected in the Application Settings. An incorrect field separator can result in all data for a given row appearing in a single spreadsheet cell. To change the field separator choose **Settings > Application** and then choose **Report** from the **Settings List**.

Graphing Data

Start by selecting the features to include in the graph. If both image channels are displayed, data for both channels will be included in the initial graph. To open a chart view, click  on the toolbar or choose **Report > Chart View**. Chart View can also be chosen on the contextual menu accessed by right clicking the image.

The controls in the bottom half of the Chart View window allow the chart to be changed interactively. The **Chart Style** can be used to switch between a line chart and a bar chart. The **Display Values** list shows the data value (average intensity, integrated intensity, etc.) graphed for each feature. The list also shows the **Channel** the data belong to and the **Color** used for identification. Existing values can

be changed using the drop-down menus in the **Field** and **Channel** columns. Color can be changed by clicking the color button in the appropriate row and choosing a new color from the color palette.



To remove a set of data values, click the row number (first column) to select the row and click **Remove**. To add a set of data values, click **Add** and then set the **Field**, **Channel**, and **Color** as desired.

The **X-Axis Properties** button can be used to change the label on the X-axis and to change the sort order of the features on the X-axis. The **Axis Label** field can be used to switch between the feature name and

ID. When **Sort** is selected, features on the X-Axis will be sorted in descending order using the selected **Field**. Features are sorted in ascending order when **Ascending** is selected.

The font size on the X-axis is set automatically and decreases as the number of data points increases. If more than 45 data points need to be plotted on the X-Axis, the labels become unreadable and are removed automatically. Information about a specific data point is displayed in a tool tip when the cursor is moved over the data point and stopped.

Using Templates

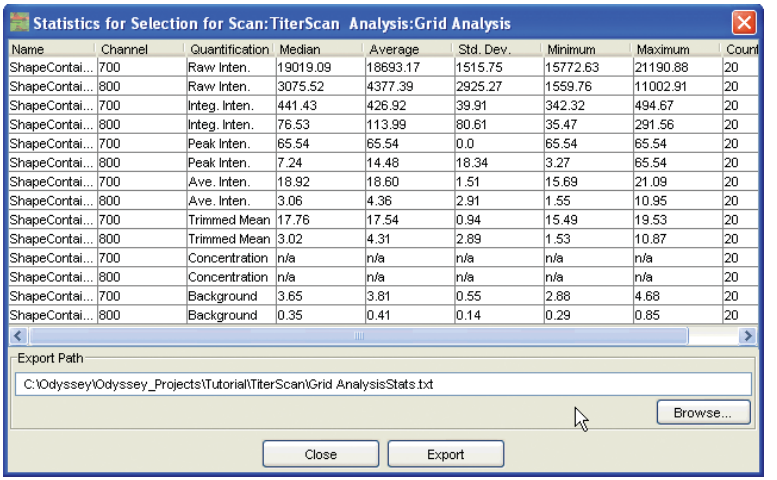
Chart View templates store the **Chart Style**, **X-Axis Properties**, and **Display Values** so they can quickly be loaded using the **Load From Template** list in the Chart View window. After a template is loaded, it can be changed by clicking **Modify**. The **Chart Style**, **X-Axis Properties**, and **Display Values** in the Edit Template window operate as described above for the Cart View window. After editing the template, click **Save** to change the current template or **Save As** to create a new template. New templates can also be created by choosing **Settings > Chart View Templates**.

Displaying and Exporting Statistics

Statistics can be displayed for a set of features by selecting the features and choosing **Report > Stat Table View**. The Statistics Table displays the median, average, standard deviation, minimum and maximum for the following variables for the set of selected features:

- Raw Integrated Intensity
- Integrated Intensity
- Peak Intensity
- Average Intensity
- Trimmed Mean
- Concentration
- Background

Each of these variables is described in the Chapter 13.



Name	Channel	Quantification	Median	Average	Std. Dev.	Minimum	Maximum	Count
ShapeContai...	700	Raw Inten.	19019.09	18693.17	1515.75	15772.63	21190.88	20
ShapeContai...	800	Raw Inten.	3075.52	4377.39	2925.27	1559.76	11002.91	20
ShapeContai...	700	Integ. Inten.	441.43	426.92	39.91	342.32	494.67	20
ShapeContai...	800	Integ. Inten.	76.53	113.99	80.61	35.47	291.56	20
ShapeContai...	700	Peak Inten.	65.54	65.54	0.0	65.54	65.54	20
ShapeContai...	800	Peak Inten.	7.24	14.48	18.34	3.27	65.54	20
ShapeContai...	700	Ave. Inten.	18.92	18.60	1.51	15.69	21.09	20
ShapeContai...	800	Ave. Inten.	3.06	4.36	2.91	1.55	10.95	20
ShapeContai...	700	Trimmed Mean	17.76	17.54	0.94	15.49	19.53	20
ShapeContai...	800	Trimmed Mean	3.02	4.31	2.89	1.53	10.87	20
ShapeContai...	700	Concentration	n/a	n/a	n/a	n/a	n/a	20
ShapeContai...	800	Concentration	n/a	n/a	n/a	n/a	n/a	20
ShapeContai...	700	Background	3.65	3.81	0.55	2.88	4.68	20
ShapeContai...	800	Background	0.35	0.41	0.14	0.29	0.85	20

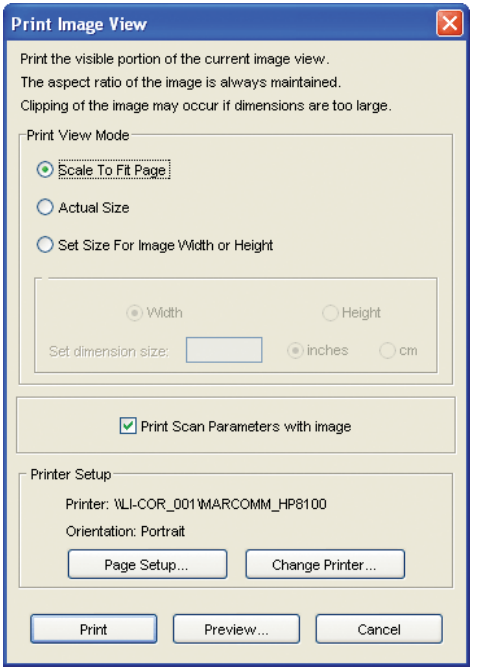
The **Count** column indicates the total number of features selected for a given channel. For the statistics shown above, 20 features were selected in both the 700 channel and 800 channel (channels were overlaid when 20 features were selected).

To export the statistics in tab-delimited text format, click **Browse** to select the path and click **Export**. (The path can also be typed in the **Export Path** field.

Printing an Image View

Images displayed in an Image View window can be printed by choosing **File > Print Image View**.

To scale the image so the printed image fits the current page size, click **Scale to Fit Page** in the Print Image View window. To print the image at actual size, click **Actual Size**. To constrain the printed image so it prints at a certain height or width, click **Set Size for Image Width or Height**, select **Width** or **Height** and enter the size to which the image should be constrained.



Scan parameters (resolution, etc.) can be printed with the image by selecting **Print Scan Parameters with Image**. Click **Page Setup** to select portrait or landscape page orientation. To change printers before printing, click **Change Printer**. The Preview button displays the image as it will appear on the printed page.

After making changes, click **Print** to send the image and all annotations to the printer. The visible portion of the image is printed with all current features including annotations, brightness and contrast changes, zoom, etc. Any portion of the image that is scrolled outside of the window boundaries will not be printed. For pseudo color images, the pseudo color legend will be appended to the right side of the image.

Note: *If the Application Settings (Image View Display) have been used to enlarge the text display area in order to prevent annotations from being truncated at the edge of the image, this extra text display area is included when an image view is printed (assuming the extra text area is visible in the Image View window).*

Viewing and Printing the Scanner Log

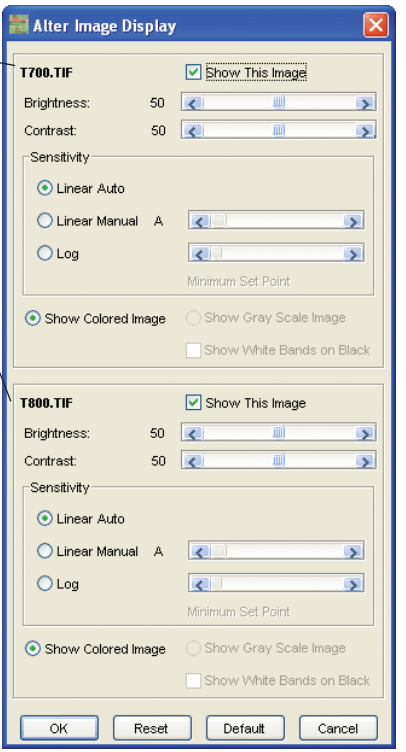
Choosing **Report > Scanner Log** displays the scanner log of the Odyssey Imager. From this window the log can also be printed. The scanner log is useful for diagnosing problems. LI-COR technical support may request a copy of the scanner log when diagnosing problems.

Chapter 11: Changing the Appearance of Scanned Images

Image Display Adjustments

Choose **View > Alter Image Display** to open the Alter Image Display window. The Alter Image Display window can also be opened by clicking  on the tool bar.

The name of the image corresponding to each group of controls is shown in the upper left corner.

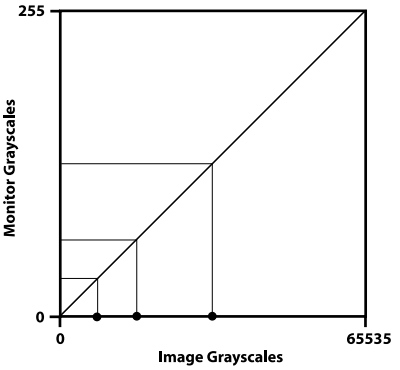


Each image has its own set of **Brightness**, **Contrast**, and **Sensitivity** sliders. The **Brightness** and **Contrast** sliders change the appearance of the background and bands in the image.

Changing How Image Data Are Mapped to the Monitor

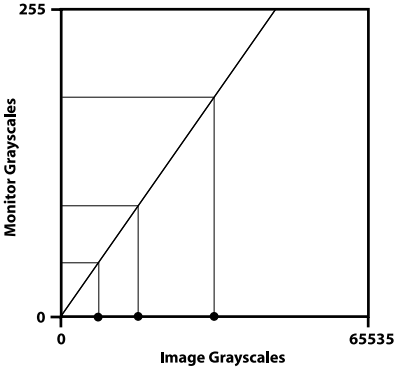
If a scan appears blank, the cause may be due to failed reactions, but it may also be because the sensitivity needs adjustment. Odyssey image files contain over 65000 grayscale values, but the typical computer monitor can display only 256 grayscale values. This requires a scheme to "map" grayscale values in the image to the monitor. Three **Sensitivity** controls are provided that give alternative methods of mapping grayscale values to the monitor. Typically, either **Linear Auto** or **Linear Manual** gives the best results. A manual **Log**arithmic-mapping method is also provided.

Linear Auto uses computer algorithms to predict the sensitivity that will give the best image appearance. When **Linear Auto** fails, choose **Linear Manual** and use the sensitivity slider to interactively change the sensitivity until the image appears the way you want. Linear mapping, which is preferred in most applications, can best be explained by considering a graph that has image grayscale values on the X-axis and monitor grayscales on the Y-axis.



Linear grayscale mapping.

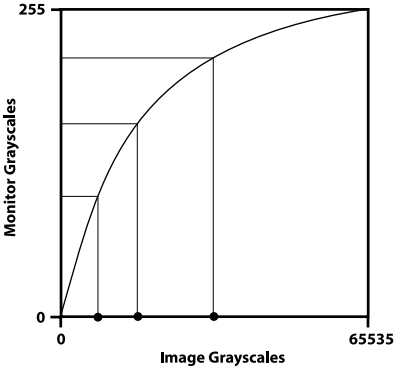
With **Linear** mapping, a given change in image intensity results in a proportional change in display intensity. The three filled circles on the X-axis above represent the grayscale values of three bands of relatively low intensity. When bands have low intensity on the monitor, the most common way to "intensify" the bands is to change the sensitivity setting. This changes the slope of the linear response line as shown below. The result is that low intensity bands are displayed with higher grayscale values on the monitor.



Linear grayscale mapping after increasing sensitivity.

Increasing the sensitivity will produce satisfactory results as long as there are no bands on the image with high grayscale values. Bands with high grayscale value will saturate as the sensitivity is increased. When there are bands over the entire grayscale range but weak

bands need to be intensified, **Log**arithmic mapping can help because it intensifies the display of weak bands while preventing strong bands from saturating.



Logarithmic mapping intensifies weak bands while keeping strong bands from saturating.

Note: *These image display controls only influence image display and do not change image data.*

Changing Image Display Style

Most analysis functions are performed with only one of the two images displayed. The **Show This Image** check boxes in the Alter Image Display window are used to show or hide one of the two images. Similar controls for selecting which image to display are discussed later in this chapter.

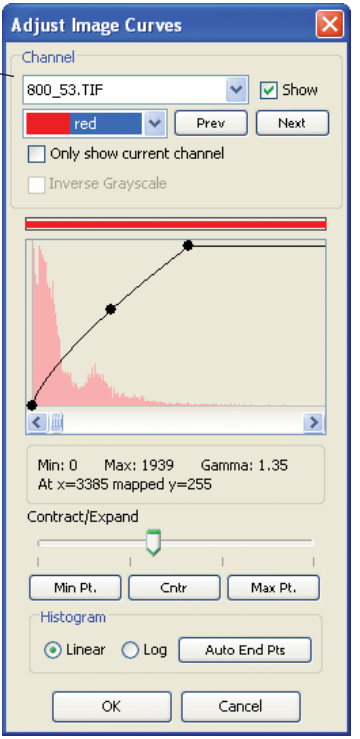
If only one image is displayed, the image can be switched between color and black-and-white using the **Show Colored Image** and **Show Gray Scale Image** radio buttons. When an image is displayed as grayscale, the **Show White Bands on Black** check box can be used to invert the image display style to white bands on a black background. (Normally fluorescence shows up as dark bands on a "white" background.)

The **Reset** button changes the **Brightness**, **Contrast**, and **Sensitivity** back to the way they were when the window was opened. The **Default** button changes the image display settings to the default programmed settings.

Adjusting Image Curves

Adjust Image Curves window can be opened by clicking  on the tool bar or choosing **View > Adjust Image Curves**. The Adjust Image Curves window can also be opened from the Scanner Console window or the New Analysis window.

The channel area is used to select the channel to view and change its appearance.



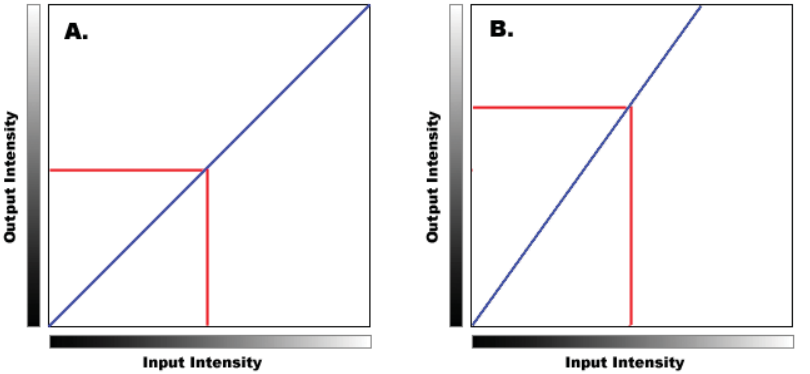
Note: *If the images were previously adjusted using the* ***Alter Image Display*** *function, a message may be displayed indicating the image curve settings are not stored for the image. There is not always a direct conversion between the two image manipulation methods used in* ***Alter Image Display*** *and* ***Adjust Image Curves****, so defaults are used when switching back and forth between the two methods.*

Start by selecting the channel (700 or 800) to adjust. The **Channel** drop down menu or the **Next** and **Prev** buttons can be used to switch channels. A histogram of intensity values and an adjustment curve for the selected channel are displayed in the middle of the window. Any adjustments made to the curves are immediately shown on the image(s) in the Image View window. The **Show** and **Only Show Current Channel** check boxes can be used to determine whether one or both image channels are displayed. The color of the current image can be changed using the color drop down menu (the colors in the Application settings for new analyses are not changed). **Pseudo Color** is available as a color if you have purchase a key to unlock the Odyssey MousePOD™ module (see *Odyssey In vivo Imaging Guide*). The **Invert Grayscale** check box inverts the color map (whites become black and blacks become white) for grayscale and pseudo color images.

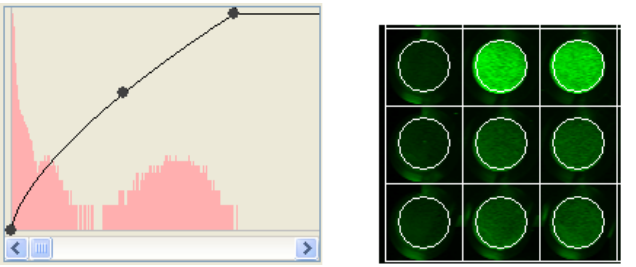
Using the Intensity Adjustment Curve

The intensity adjustment curve graphically illustrates how grayscale values in the image file are mapped to the display. As indicated below, the X-axis is the original or input intensity value from the TIFF image file. The Y-axis is the grayscale value that is output to the display. For both axes, dark pixels (lowest intensity) are located near the origin. The intensity adjustment curve is used to calculate the output intensity for a given input intensity.

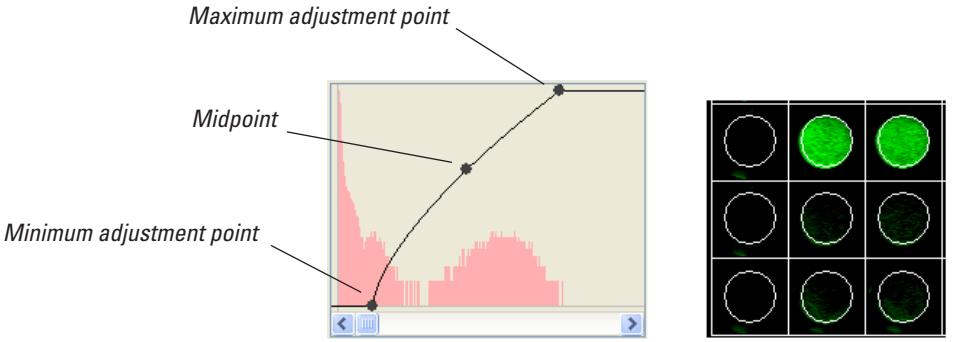
For purposes of illustration, a linear intensity adjustment curve is shown in figures A and B below.



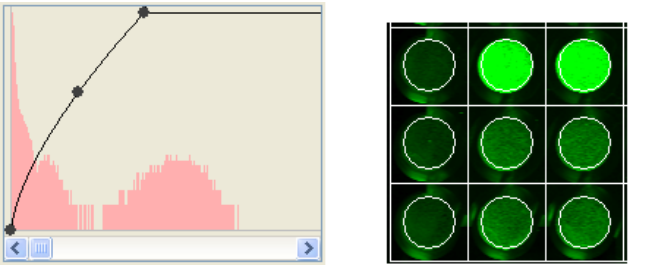
Note that as the slope of the curve increases (B), the output intensity of a given pixel increases. The adjustment curve shown in the Adjust Image Curves window has three adjustable points that appear as dots on the curve. These three points can be used to change the slope and shape of the curve. The initial shape of the adjustment curve and location of the three adjustment points are calculated by Odyssey software. To revert to the initial calculated curve, click **Auto End Pts**.



Above is an unedited curve and the corresponding image data from the 800-channel. Compare these data to the curves and images below (the illustrations are shown in color in the Odyssey help system).

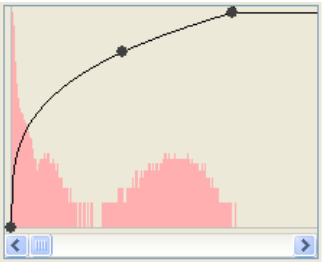


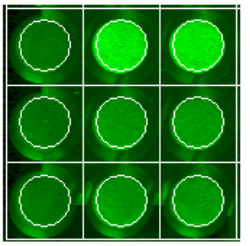
The adjustment curve and image above illustrate several effects of moving the minimum adjustment point (lower left) toward the center (it only moves horizontally). First, many of the darker pixels in the input file are now mapped to black on the monitor. (All pixels to the left of the adjustment point are mapped to black.) The result is that areas composed primarily of dark pixels are now nearly black. The second effect is a slight increase in the slope of the adjustment curve, which will make some of the brighter pixels even brighter.



The adjustment points have additional functions with pseudo color images. See *Odyssey In vivo Imaging Guide* for details.

The maximum adjustment point (upper right) can also be moved horizontally. As the adjustment point is moved to the left, all pixels to the right of the adjustment point are mapped to maximum intensity and become much brighter. As the maximum adjustment point is moved to the left, the slope of the adjustment line increases, making many of the other pixels brighter as well.

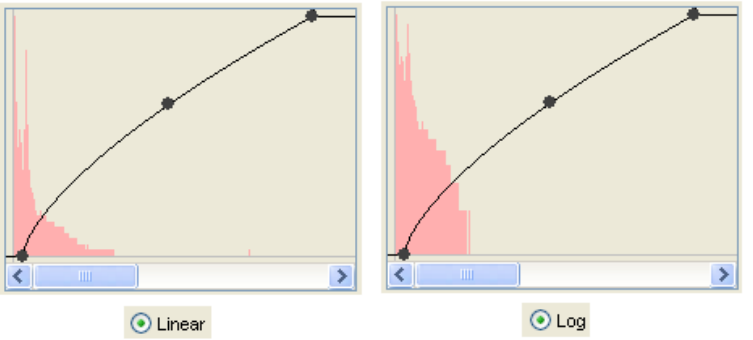




The midpoint can be dragged in only the vertical dimension. As the midpoint is moved, the shape of the adjustment curve is changed, which results in dark pixels and bright pixels being adjusted by different amounts. It general, changing the midpoint will have the greatest impact on the mid tone pixels, but the bright and dark pixels will also be changed.

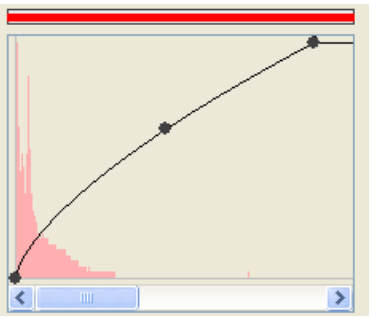
Using the Histogram

Linear vs. Log: The histogram shows the number of pixels (Y-axis) that have a given grayscale intensity value (X-axis). When **Linear** is selected in the Adjust Image Curves window, the histogram is plotted with a linear Y-axis. For images that have many dark pixels and comparatively few pixels with high intensity values, a linear Y-axis can make it difficult to see high intensity values. Clicking **Log** to switch to a log scale for the Y-axis can make it easier to see intensity values with fewer pixels.



The X-axis of the histogram can be expanded or contracted using the **Contract/Expand** slider. The scroll bar underneath the histogram can be used to scroll along the X-axis. When the X-axis is expanded, the **Min Pt.**, **Cntr**, and **Max Pt.** buttons can be used to quickly scroll to the minimum point, midpoint (center), and maximum point, respectively.

The bar over the top of the histogram indicates the maximum intensity. If the cursor is stopped over the bar, the maximum intensity is listed in the tool tip. If maximum intensity its very low, the image may have been scanned with the intensity parameter set too low to get adequate signal strength. Conversely, if intensity is set too high during scanning, saturated pixels may result. A red bar all the way across the intensity indicator, as shown below, indicates saturation.



Cropping, Rotating, and Flipping Images

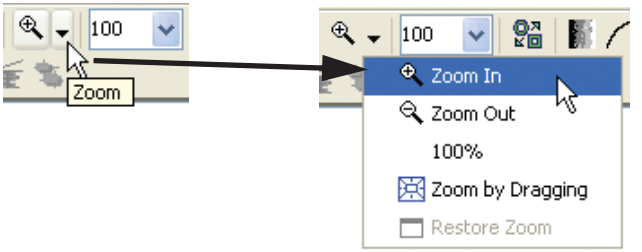
In Odyssey, most image manipulations like cropping, flipping, and rotating are done in the New Analysis window as described in Chapter 3. When a new analysis is added, the image can be flipped, rotated, cropped, or filtered as needed. The background fluorescence can also be subtracted. These functions are accessible only when a new analysis is created.

Magnifying the Image

All of the zoom functions described in this section can also be accessed via the **View** menu.

Zoom Functions on the Toolbar

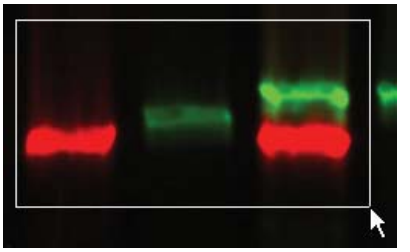
Click  on the toolbar to magnify the image. Click the down arrow next to the magnify icon to access other functions.



Zoom Menu Choices:

- **Zoom In**: Same as described above
- **Zoom Out**: Decreases image magnification

- **100%**: Returns the zoom level to 100%
- **Zoom by Dragging**: Click and drag a rectangle around an area to zoom in on and release the mouse button when the area is enclosed.



The rectangular region is enlarged to fit in the current Image View window. To get a lot of magnification, enlarge the Image View window before using the **Zoom by Dragging** function.

- **Restore Zoom**: **Restore Zoom** works in conjunction with **Zoom by Dragging** and resets to the zoom level prior to using **Zoom by Dragging**.

Use the drop-down magnification list to select a specific magnification (percent of original size).

Keyboard Shortcuts

Function key **F11** increases image magnification. Unlike **Zoom In** on the **View** menu, however, the zoom is centered at the cursor position as long as the cursor is over some part of the image. Function key **F12** zooms out, centered at the cursor position.

Overlaid Images

For scans that have both 700- and 800-channel images, the images are automatically overlaid when an analysis is opened. To display a single image, choose **View > Single Channel** or click the overlay button ( ) on the tool bar. The overlay button toggles between **Channels Overlaid** and **Single Channel** display.

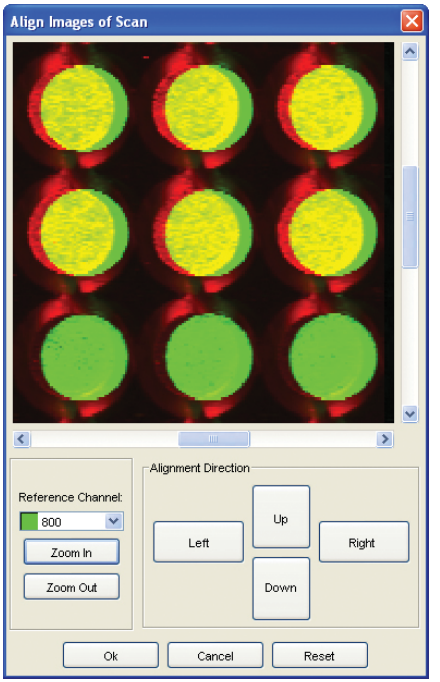
The **View** menu changes when a single channel is displayed. The **Single Channel** menu changes to **Channels Overlaid**, which is used to overlay the images again. In addition, menus are activated to switch between channels and to switch between various image display styles.

Icons on the tool bar are activated to change the image display style from color to grayscale ( ), or back to color ( ), as well as pseudo color ( ) if you have the optional Odyssey MousePOD™ Module. To switch between channels, click ( ).

Aligning Images

When scanning at high resolution (84, 42, and 21 µm), it is possible for the 700- and 800-channel images to be misaligned. The image below shows a microplate scan where the 700-channel image (red) is horizontally offset from the 800-channel (green) image. Images can be aligned by choosing **File > Align Images**.

Important: There is no undo operation for image alignment because the TIFF image files are immediately changed.



To correct an offset, start by using the zoom buttons to make the offset clearly visible. Next, select the reference channel that stays stationary while the other channel is moved. Last, move the channel not designated as the reference using the **Alignment Direction** buttons. The **Up**, **Down**, **Left**, and **Right** buttons move the image one pixel at a time. To start over, click **Reset**.

Note: *The Application settings can be used to enable automatic image alignment when the Adjust Locations function is used to move features (see Chapter 7).*

Changing to Grayscale Image Display Style

If a single image is displayed (rather than two images overlaid), the image display style can be switched to grayscale by clicking  in the tool bar or choosing **View > Grayscale**.

The Alter Image Display window can also be used to switch between grayscale and color display as discussed earlier in this chapter.

Changing to Color Image Display Style

While a single image is displayed, the image display style can be switched to color by clicking  in the tool bar or choosing **View > Color**. The Alter Image Display window can also be used to switch to the color display style as discussed earlier in this chapter.

Changing to Pseudo Color Image Display Style

The pseudo color display tool on the tool bar is grayed out unless you have purchased a key to unlock the Odyssey MousePOD™ Module. See the *Odyssey In vivo Imaging Guide* for details on pseudo image display.

Switching Between Image Channels

If a single image is displayed (rather than two images overlaid), clicking  in the tool bar or choosing the channel on the **View** menu switches between the 700-channel image and the 800-channel image. If the 700-channel image is displayed, **Channel 800**

can be chosen from the **View** menu as shown below. If the 800-channel image is displayed, the last choice on the **View** menu will be **Channel 700**.

Displaying a Second Image View Window

For scans with both a 700- and 800-channel image, a second image view can be opened by choosing **View > Display 2nd View** or by clicking  on the tool bar.

The two image view windows are displayed side-by-side in the Odyssey window. If the first image view was in single channel mode before selecting **Display 2nd Channel**, the second image view will be the opposite channel. If the first image view shows both channels overlaid, the second image view will be the 800-channel image in single channel mode.

Hiding Image Annotations

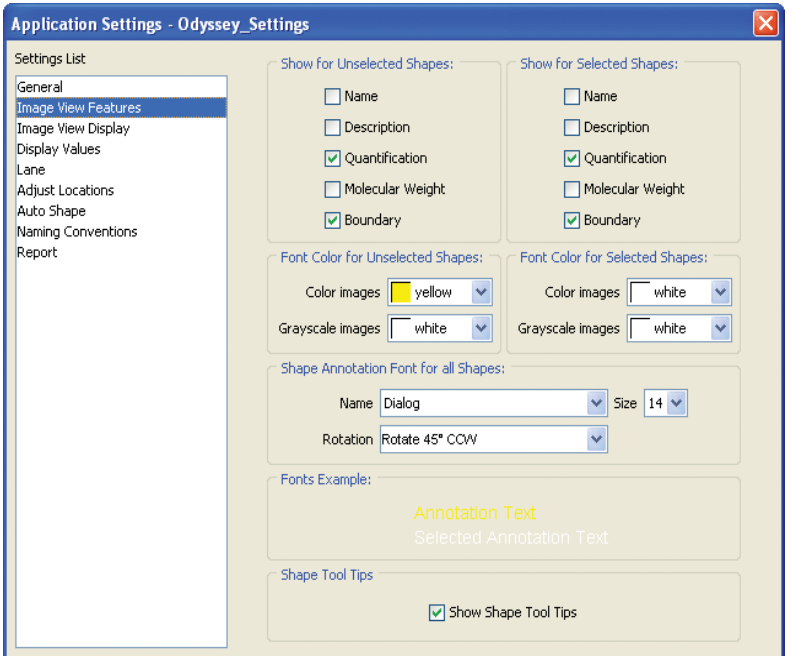
If annotations obscure other image features, all annotations can be temporarily hidden by clicking  on the tool bar or choosing **View > Hide Annotations**.

To view hidden annotations, click  on the tool bar again or choose **View > Show Annotations**.

To reduce screen clutter, the Application settings can be used to control which annotations are displayed, as described below.

Using the Application Settings to Display Labels

Image View Feature settings are opened by clicking  on the tool bar or by choosing **Settings > Application** and then selecting **Image View Features** from the settings list. Image View Feature settings are important when analyzing membranes because they can be used to display integrated intensity, molecular weight, or other values over each band/spot. For grids, labels are not displayed regardless of how the Image View Features are set. (Manually added text annotations are displayed on grids.)



The Image View Feature settings can be used to reduce the "screen clutter" that can occur when labels are displayed on closely spaced features. Notice that there are two groups of settings – those for

selected features and those for *unselected* features. One strategy to reduce screen clutter is to treat selected and unselected features differently. If all desired labels for selected features are turned on and all labels except boundaries are turned off for unselected features, the display will stay uncluttered and individual features can be clicked to display their labels.

In other situations, like exporting annotated images, it is useful to have labels for unselected features turned on. Turning on quantification values for unselected features, for example, displays all quantification values on the image.

Each type of label is described below:

- **Name:** Name is either the auto-entered default name or a name entered in the properties.
- **Description:** Description is blank by default for many features, but can be entered in the properties.
- **Quantification:** Quantification values are displayed as Integrated Intensity or Concentration depending on the Application settings.
- **Molecular Weight:** The Molecular Weight annotation displays band migration in the gel as molecular weight, scan line on the image, or percentage of the lane according to the Application settings (described below). Molecular weight values are displayed as "n/a" (not assigned) until molecular weight standards have been identified.
- **Boundary:** Each feature (circle, etc.) has a boundary. If boundaries are turned off only the text annotations, if any, will remain.

Changing Font Specifications

The font used to annotate features can also be changed to reduce screen clutter. Changing the normal or selected color of the font, reducing the size of the font, or even changing to a different font can improve the readability of annotations on the image. A sample of the currently selected font and font colors is shown in the **Fonts Example** area at the bottom of the window. Separate controls are provided to

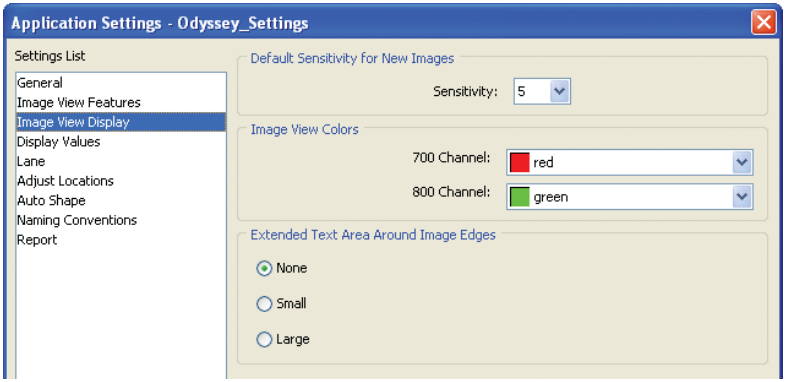
pick colors for both color and black-and-white image display styles (some colors do not work well for both display styles). Annotations can be rotated 45 or 90 degrees counter-clockwise using the **Rotation** drop-down list.

Displaying Data in Tool Tips

When the **Show Shape Tool Tips** check box is selected, data values are displayed in a tool tip when the cursor is stopped over a feature.

Using the Image View Display Settings

The Image View Display settings are opened by choosing **Settings > Application** and selecting **Image View Display** from the **Settings List**.



Setting the Default Sensitivity for New Images

When a new scan is opened for the first time, the image is displayed using the sensitivity value shown in the Image View Display settings. Generally, the **Auto** setting provides good results. However, if the

sensitivity always needs adjustment after opening new images, the default sensitivity can be changed by picking a value from the **Sensitivity** list.

Changing Image Colors From Red/Green

Using the **700 Channel** and **800 Channel** drop-down lists, the color of the 700- and 800-channel images for any new analysis can be changed to red, green, or blue. Each channel must have a different color to distinguish fluorescence from each channel when channels are overlaid. Choose **View > Adjust Image Curves** to change colors for the current analysis.

Extending the Text Display Area Around Images

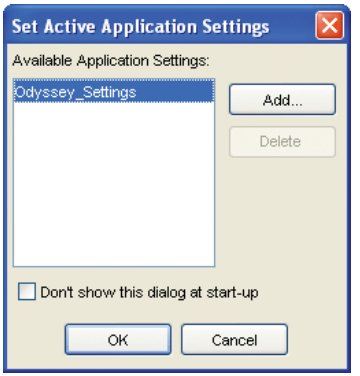
When features or annotations are drawn near the edge of an image, some of the text (data values, etc.) associated with the feature may be truncated at the edge of the image. This is most likely to happen when **Extended Text Area Around Image Edges** is set to **None** (the default). If some of the text on an image is truncated, increase the text display area by setting **Extended Text Area Around Image Edges** to **Small** or **Large**. This does not add pixels to the image data, it only allows truncated text to be displayed.

Note: *This additional text display area is included when an image view is printed (**File > Print Image View**) or exported to a file (**File > Export Image > Export Image View**).*

Chapter 12: User Accounts and Settings

Application Settings

When the Odyssey application starts, the window shown below is displayed, which can be used to choose the application settings for the current session. This makes it easy for users to have their own settings file that determines important parameters such as background calculation method. In labs that do not have multiple users, however, the settings selection window is not needed. When **Don't Show This Dialog At Startup** is selected, the settings selection window will no longer be displayed when the Odyssey application starts. The active settings file can be changed at any time by choosing **Settings > Select Active Settings**.



To add a new set of settings, click **Add** and name the settings file. A new settings file will be created containing default settings.

A settings file can also be created by copying the active settings file. To copy a settings file, choose **Settings > Select Active Settings**, select an application settings file and click **OK**. Next, choose **Settings > Application**, change any of the application settings, and save a new settings file by clicking the **Save As** button and naming the new file.

Note: *Only Application settings are stored. Other settings such as ICW Setup are not stored.*

To delete a settings file, select it in the Set Active Application Settings window, and click **Delete**.

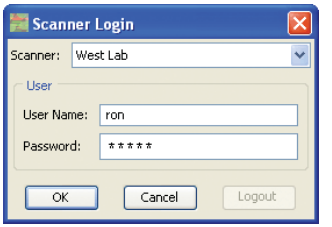
Note: *The last settings file cannot be deleted.*

User Administration

The **User Administration** menu choice on the **Settings** menu is used to change your own password and scan groups. User accounts, passwords, and scan groups are stored on the Odyssey instrument. See *System Administration* below to make changes to user accounts or scan groups other than your own.

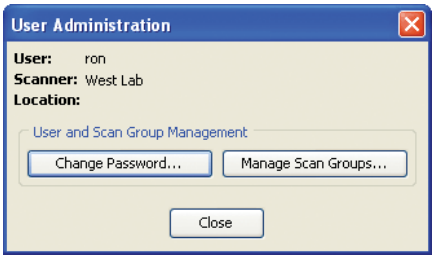
Changing Your Own Password

1) Choose **Settings > User Administration**.



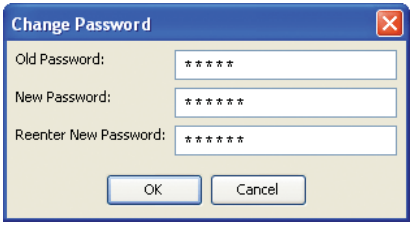
*If someone else is already logged in, click **Logout** before entering your **User Name** and **Password**.*

2) Select a scanner from the **Scanner** list and enter the **User Name** and **Password** of your account. Click **OK** to continue. (See Scanner Settings if no scanners are listed.)



3) Click **Change Password** and enter your current password in the **Old Password** field.

4) Enter a new password in the **New Password** field and repeat it in the **Re-enter Password** field.



5) Click **OK** to change your password.

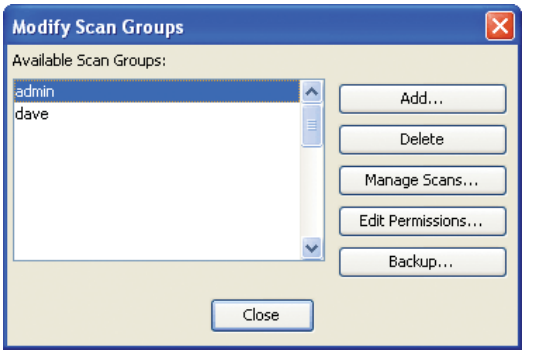
Managing Your Own Scan Groups

1) Choose **Settings > User Administration**.

2) Select a scanner from the **Scanner** list and enter the user name and password of your account. Click **OK** to continue.

3) Click **Manage Scan Groups**.

Scans are stored on the Odyssey instrument in scan groups. Scan groups are like folders that require permission to access them. Normally, your own scan group (matches your user name) is listed, along with any other scan groups that may have been added.



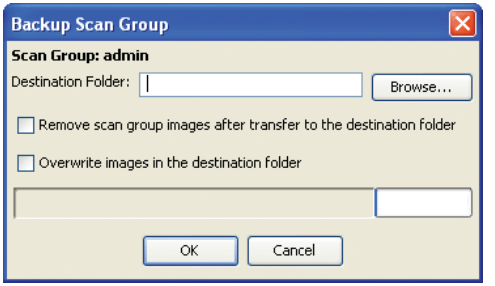
The following tasks can be accomplished in the Modify Scan Groups window:

- **Add Scan Groups:** Click **Add** and enter the name of a new scan group. The user who created the scan group is automatically given Control access permission. All other users have no access unless their permisssion level is changed, as described below.
- **Delete Scan Groups:** To delete a scan group on the instrument, a user with Control rights can select any of the scan groups in the Modify Scan Groups window and click **Delete**.

  IMPORTANT: *Deleting a scan group deletes all images within the group. After a group is deleted, it is not possible to restore images that were in the group. Images stored in other scan groups on the instrument or on the computer are not deleted.*
- **Delete Scans In A Scan Group:** Deleting a scan from a scan group on the instrument does not delete any scans that may be stored on the computer. To delete a scan from a scan group, select the scan group in the Modify Scan Groups window and click **Manage Scans**. All scans in the scan group are listed. Select a scan and

click **Delete** to permanently delete a scan and all associated files (image files, scan logs, etc). To delete multiple scans, use **Shift+Click** or **Ctrl+Click** to select additional files before clicking **Delete**.

- **Edit Permissions To Your Scan Group:** Any user with Control permission can allow other users to access their scan group to start new scans, analyze scans or delete scans that in the scan group. To change access for a given user, select the scan group in the Modify Scan Groups window and click **Edit Permissions**. All users are listed along with their current permission levels. Select a user and click the down-arrow button in the **Permission** column to drop down a list of permission levels. Select **None** to prevent a user from accessing your scan group. Select **Access** to give a user the ability to download and analyze scans in your scan group. Select **Change** to give a user the the ability to start new scans that are stored in your scan group as well as all the rights that come with **Access** permission.
- **Backup:** To quickly backup all files in a given scan group on the Odyssey instrument, users with Control permission can click **Backup** in the Modify Scan Groups window.



In the Backup Scan Group window, enter a destination folder or click **Browse** to open a standard file selection window. To move scans to the destination folder rather than copying them, select the check box labeled **Remove Scan Group Images After Transfer To**

Destination Folder. To overwrite scans that are already in the destination folder, select the check box labeled **Overwrite Images In The Destination Folder**. After configuring the backup, click **OK** to transfer the files.

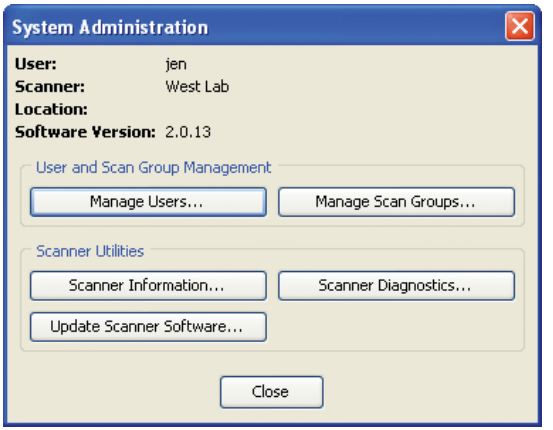
System Administration

Account Rights

To use the **System Administration** functions on the **Settings** menu, you must be logged in with an account that has Administrator access rights. There are three levels of access rights:

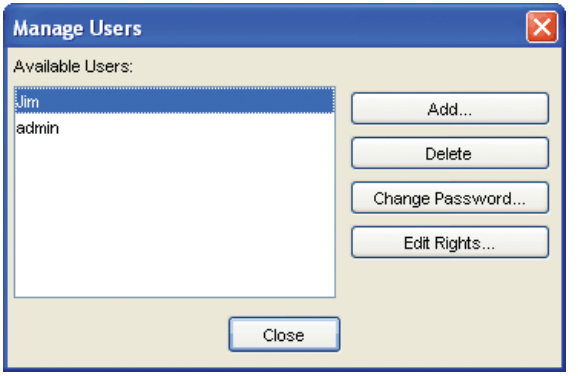
- **None.** Accounts with access rights set to None are inactive and the user cannot access data or start runs.
- **Control.** Most users are given Control access rights, allowing them to start new runs and access data in scan groups to which the user has been given access permission.
- **Administrator.** Includes all the Control rights, plus the ability to change any user account or scan group, update system software and other setup and diagnostic functions. For greatest security, accounts with Administrator access rights should not be used for daily operation.

To use the system administration functions, choose **Settings > System Administration** and log in using an account with Administrator rights.



Managing Users

Click **Manage Users** in the System Administration settings window to make changes to any user account.



Adding Users: Click **Add** in the Manage Users window.

Enter the name and password for the new user and repeat the password. Select the access rights according to the definitions given above. Click **OK** to add the user. When a user is created, a scan group in the Odyssey instrument is also created. The new scan group has the same name as the user name.

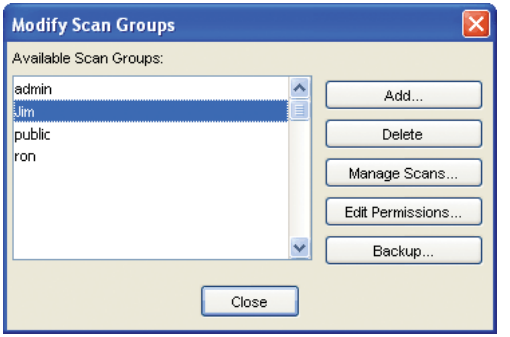
Deleting Users: To delete a user, select the user from the **Available User** list (Manage Users window) and click **Delete**. Note, however, that this only deletes the user account. Any data must be deleted by deleting the scan group using the **Manage Scan Groups** button in the System Administration window.

Changing Passwords: Passwords for any user can be changed by selecting the user from the **Available User** list in the Manage Users window and clicking **Change Password**. Enter the new password and then verify the password by entering it again and click **OK**.

Editing Access Rights: Access rights for any user can be changed by selecting the user in the Manage Users window and clicking **Edit Rights**. Click the down-arrow button in the **Permission** column next to the appropriate user to drop down a list of access rights. Choose the access rights and click **Close**.

Managing Scan Groups

To add or make changes to scan groups in the Odyssey instrument, click **Manage Scan Groups** in the System Administration window.



Adding Scan Groups: Click **Add** in the Modify Scan Groups window, enter a name for the group, and click **OK**.

Editing Access Permission to a Scan Group: Select the scan group to edit and click **Edit Permissions**. Click the button in the **Permission** column across from the user name whose access permission you want to change. Select a new permission level and click **Close**. Select **None** to prevent a user from accessing the scan group. Select Access to give a user the ability to download and analyze scans. Select Control to give a user the ability to start new scans that will be stored in the scan group as well as all the rights that come with Access permission.

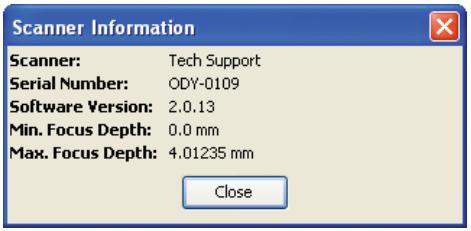
Deleting Scan Groups: Select a scan group from the **Available Scan Groups** list and click **Delete**. The scan group and any scans stored in it are deleted.

Deleting Scans: Select the scan group containing the scan to be deleted from the **Available Scan Groups** list and click **Manage Scans**. Select the scan from the list of scans and click **Delete**.

Backup: To quickly backup all the files in a given scan group from the Odyssey instrument to the computer, click **Backup** in the Modify Scan Groups window. In the Backup Scan Group window, enter a destination folder or click **Browse** to open a standard file selection window. To move the scans to the destination folder rather than copying them, select the check box labeled **Remove Scan Group Images After Transfer To Destination Folder**. To overwrite scans that are already in the destination folder, select the check box labeled **Overwrite Images In The Destination Folder**. After configuring the backup, click **OK** to transfer the files.

Scanner Information

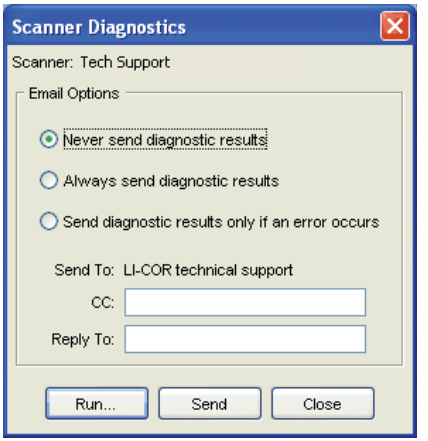
Click **Scanner Information** in the System Administration settings to display the Scanner Information window. The items displayed include the scanner name, serial number and software version number of the instrument software. The minimum and maximum focus depth are the smallest and largest values that will be accepted for the focus offset parameter when starting a scan.



Scanner Diagnostics

The Odyssey instrument has a built-in procedure to diagnose hardware problems that could result in faulty operation. If you are experiencing difficulty with your instrument, the first step is to contact LI-COR Technical Support. Someone in Technical Support may ask you to run the scanner diagnostic routine. This routine runs

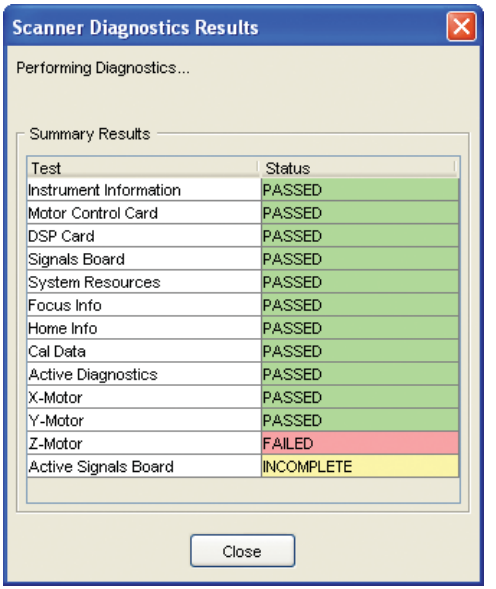
a series of tests on various hardware components. The results can be e-mailed to LI-COR Technical support automatically if your network permits such operations.



Start by selecting one of the three options for e-mailing the results:

- **Never Send Diagnostic Results**: Displays results but does not e-mail them.
- **Always Send Diagnostic Results**: Always sends results to LI-COR Technical Support and also to any e-mail address included in the carbon copy (**CC**) field. The **Reply To** field should be used to include the address of the sender.
- **Send Diagnostic Results Only If An Error Occurs**: Same as above except that results are e-mailed to LI-COR only if one of the diagnostic tests fails.

Click **Run** to start the diagnostic test. Normally, all tests will have a status of PASSED if the hardware is operating normally. A status of FAILED indicates a problem and INCOMPLETE may also indicate a problem.



Scanner Update

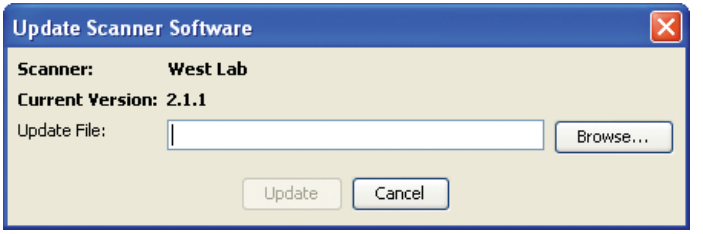
Updates to the Odyssey instrument software will be posted on LI-COR's FTP site and made available on CD-ROM. Update notifications will be sent that explain how to get the update on CD and will also include the FTP address, name, and password to download the update file.

IMPORTANT: The update software temporarily uses space on the Odyssey hard disk while the update is running. If the hard disk is too full, the oldest scans will be deleted as needed. Make sure all scans have been downloaded from the Odyssey before proceeding.

Whether you have an update CD or have downloaded the update files via FTP, the update procedure is as follows:

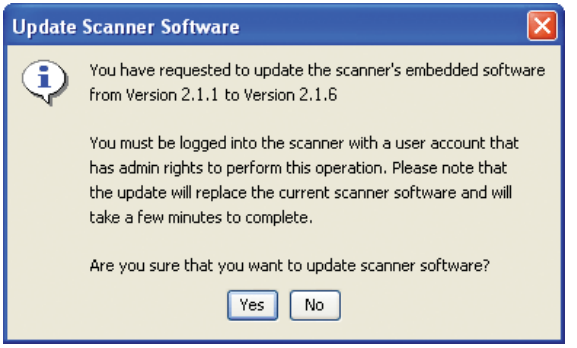
1) If you have an update CD, insert the CD into the drive. If you downloaded the update file, make sure the file is accessible on your computer or network drive.

2) Click **Update Scanner Software** in the System Administration window to open the Scanner Update window.



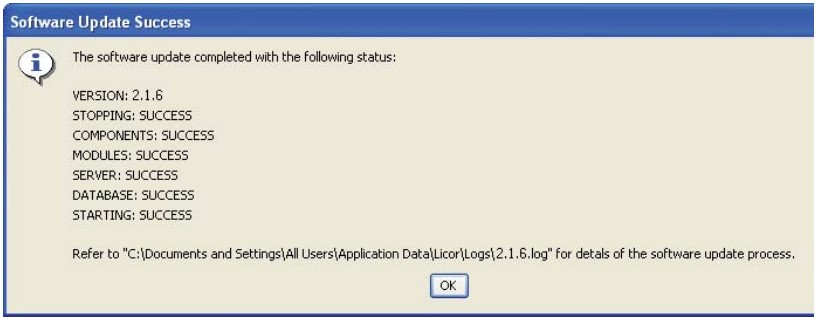
3) Click **Browse** and use the standard file selection window to select the update file from CD or disk drive. If the update file name is not displayed, make sure that all file types are being displayed in the file selection window.

4) Click **Update**. An information window is displayed that explains what the update procedure is going to do.



5) Click **Yes** to start the software installation.

There will be several minutes of apparent inactivity while the new software is being uploaded. When the update is complete, a report is displayed that indicates the success of the update procedure. If the update was successful, the word “success” is displayed next to each update component as shown below. Contact LI-COR Technical Support if any of the update components fail. Technical Support may request a copy of the update log. The path to the update log file (a text file) is displayed at the bottom of the update report window.

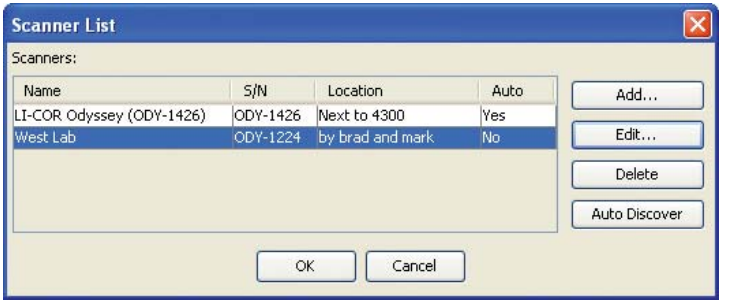


Adding and Deleting Scanners

The network address of any Odyssey instrument (scanner) must be added to Odyssey software before the instrument can be operated. Odyssey instruments are generally added during installation, but this section describes using the Scanner settings in the event of network changes, or if the instrument is moved to a different network.

Each time Odyssey software starts, any scanners on the same network as the computer are automatically detected and added to the list of available scanners. If the scanner is on a different network than the computer and the networks are connected by a router, Odyssey software will not automatically detect the scanner, but the scanner can be added to the scanner list manually. (Chapter 4 of the Odyssey

Operator's Manual has complete details on network setup for the Odyssey instrument.) To add, delete, or edit a scanner on the scanner list, choose **Settings > Scanners**. The Scanner List contains all Odyssey instruments that have been automatically detected or manually added to Odyssey software.



Adding Scanners via Auto Discovery

When **Settings > Scanners** is chosen, any scanners detected when Odyssey software was started are shown in the scanner list. If a scanner is turned on or added to the same network as the computer, the scanner list can be updated by clicking **Auto Discover** to start the automatic discovery routine. If an instrument is not auto-discovered, it has to be added manually as described below.

Note: *If the computer and Odyssey instrument are on different networks that are connected by a router, the Odyssey instrument will not be found during automatic discovery.*

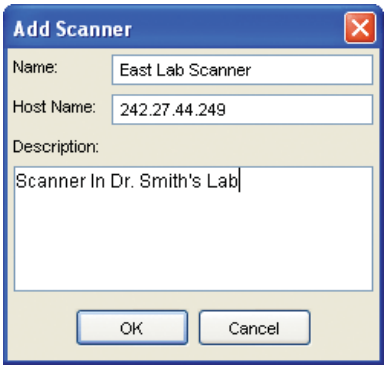
Manually Adding Scanners

Use the following steps to manually add a scanner to the list of scanners.

1) Click the **Add** button in the Scanner List dialog.

2) Enter a **Name** that identifies the scanner.

3) Enter a **Host Name**. The Host Name must be the IP address of the Odyssey instrument or a valid host name. If a host name is used, it must be registered with a Domain Name Server (DNS) available to the network.



*Note that leading zeros are not entered in the **Host Name** field. For example, you may be told that the IP address is 242.027.044.249 (a fictitious address), but the address should be entered as 242.27.44.249 as shown in the Add Scanner window to the left.*

4) Enter a **Description** to identify the scanner (optional).

5) Click **OK** to add the scanner.

Editing and Deleting Scanners

Scanners can be edited or deleted by opening the Scanner Settings, selecting the name in the scanner list, and clicking the **Edit** or **Delete** button. Editing a scanner may be necessary if the IP address for the scanner has changed, which can occur when using automatic addressing (DHCP).

Odyssey software does not automatically delete scanners from the scanner list because some scanners may be turned off or on another networks where they cannot be automatically detected. If a scanner has been removed from your network, a scanner can be deleted from the scanner list by selecting it and clicking the **Delete** button.

Chapter 13: Calculation Descriptions

One of the most important features that Odyssey brings to protein analysis on membranes is the ability to provide linear quantitative data. This chapter describes the calculations that are vital to understanding how to use the quantification tools in Odyssey software.

Derivation of the Mathematical Expressions

Understanding the derivations and assumptions supporting the mathematical expressions will make it easier to use the computed data properly.

Definitions of terms

- Feature – any area enclosed by a user using Odyssey shape tools.
- Spot – an area with increased intensity (spot) within a feature.
- Pixel – the smallest area unit of an image that is measured with a single intensity value.
- Pixel area (a, mm^2) – the physical area of an image pixel in mm^2.
- Signal intensity (or just intensity) – signal counts measured in a single pixel per unit time.
- Sample – The physical material on a membrane that generates the signal giving rise to a spot.

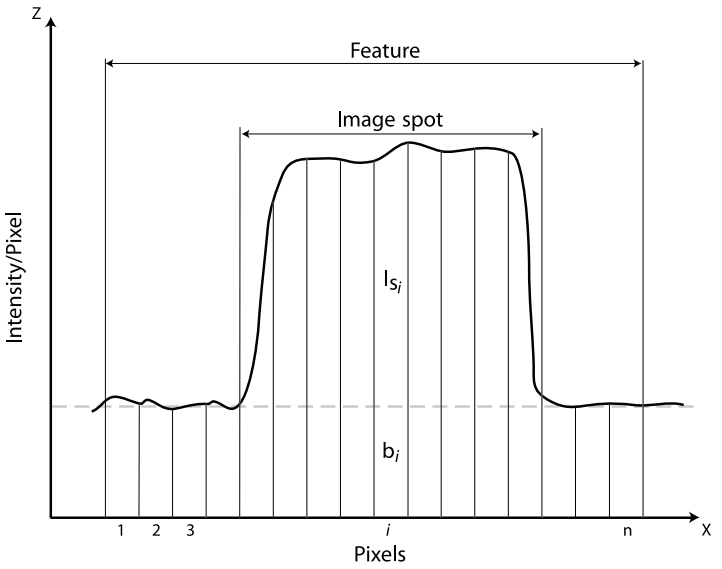
Assumptions

1) Background is uniform (though not necessarily constant) across a feature whether a spot is present or not, and regardless of the size of the feature or image spot.

2) After correcting for background, the signal in an image spot is proportional to the amount of sample generating the spot.

3) The total intensity per pixel (I) is equal to the signal intensity arising from the sample in the pixel area (I_s) plus the signal arising from the background of the pixel area (b). So, for pixel 'i',

$$I_i = I_{si} + b_i$$



Integrated Intensity and Integrated Pixel Volume

An image can be considered a three dimensional object in which two dimensions are the x and y plane of the image and the third

dimension is the pixel intensity. Consider an image containing spots with features drawn around them. The graph above shows a slice through one such feature containing an image spot. The x-axis represents pixels and the z-axis is signal intensity per pixel. This graph is a two dimensional representation of a three dimensional object that also extends along a y-axis into and out of the paper.

The total signal measured in pixel 'i' is the area (a) of the pixel times its height (I). This is called pixel volume (v). So for pixel 'i'

$$v_i = a\, I_i$$

Pixel volume is the appropriate measure of signal strength because it takes into account both the magnitude of the signal and the area over which it is distributed, which in turn is related to the distribution of sample that is generating the signal.

The total signal from the entire spot is just the summation of all the pixels and is called total pixel volume (V).

$$V = \sum_{i=1}^{n} v_i = a \sum_{i=1}^{n} I_i \qquad (1)$$

From assumption 3, $\sum I_i = \sum(I_{si} + b_i)$, and rearranging,

$$\sum I_{si} = \sum I_i - \sum b_i \qquad (2)$$

The expression $\sum I_{si}$ gives the summed intensity over the spot that is contributed by the sample after correcting for background. The last term on the right can be rewritten as $n[(1/n)\sum b_i]$, or just $n\,\bar{b}$, so $\sum I_{si} = \sum I_i - n\,\bar{b}$. Clearly, the net pixel volume of the spot V_s is just $a\sum I_{si}$, so

$$V_s = a\,(\sum I_i - n\,\bar{b}) \qquad (3)$$

The quantity calculated by equation 3 is also called the Integrated Intensity, and by assumption 2, it is proportional to the amount of sample on the membrane.

Note that the Integrated Intensity is the net pixel volume for the spot alone and is independent of feature size. If a feature is redrawn so it includes increasing numbers of background pixels, the sum of the pixels ($\sum I_i$) will increase, but since the background is assumed uniform, the increase in $n\,\bar{b}$ will compensate appropriately.

In practice, the background correction is not exact both because a uniform background is not necessarily constant and statistical fluctuations can and do occur, but also because the background is not always uniform. Nevertheless, you can confirm that Integrated Intensity is substantially independent of feature size by drawing features of varying sizes around a spot with a uniform background and observing in the Details View that the Integrated Intensity changes very little.

Note also that Integrated Intensity is substantially independent of resolution. Lower resolution means fewer and larger pixels. For features that are large compared to pixel size, the intensities recorded for pixels in similar locations over the feature will be about the same regardless of pixel size because the intensities are determined by the amount of sample and/or background at each location; but as the pixels get larger, the number of terms in the summation gets smaller, so the sum gets smaller. This is compensated by the increase in pixel area a.

Normally, background pixels are taken from three or so rows of pixels adjacent to the perimeter of a rectangle that encloses the feature drawn on the image. If one of the sides of the rectangle includes an anomaly, then either the two horizontal sides or the two vertical sides can be selected. If spots are too close to reliably isolate uniform background, then another region on the gel can be selected from which to compute $\bar{b}$; however, the number of pixels n is always

equal to the number of pixels included in the feature itself. If you want to remove background correction as a variable, set background equal to zero. In that case, it is best to draw your features so they fit the spots as closely as possible and use Integrated Intensity as the measure.

Integrated Intensity should be used as the basis for quantitative measurements.

Odyssey Calculations

Number of Pixels, Pixel Area, and Shape Area

The number of image pixels enclosed by a feature is the variable *Pixels*, which can be listed in reports. The area enclosed by a feature (mm^2) is reported in the variable *Shape Area*.

$$Shape\ Area = Pixels \times a$$

where area per pixel, a = (resolution $\times 10^{-3})^2$.

Background

Background ($\bar{b}$) is the average intensity of pixels selected as the background region,

$$\bar{b} = \Sigma b_i / n_b$$

where b_i is the intensity of the i^{th} background pixel and n_b is the number of pixels in the region selected as background.

The pixels used to calculate background are assigned by choosing **Analyze > Background Method**. The background can be set to **No Background**, **Average**, **Median**, **User Defined**, or **Lane Background Method For Bands**. Chapter 8 describes each of the background methods in detail.

Raw Integrated Intensity

Raw Integrated Intensity is defined as

$$\text{Raw Integrated Intensity} = \Sigma I_i$$

where I_i is the total intensity of the i^{th} pixel enclosed by the feature. Units are counts.

Raw Integrated Intensity has all of the properties of Integrated Intensity, except it varies with resolution. When pixel size is small compared to feature size, the values of large pixels will be similar to that of small pixels over the same area, but the number of small pixels will greatly exceed the number of large pixels so the sums will also vary. Raw Integrated Intensity is not multiplied by pixel size to correct for this. Therefore, it makes comparisons between experiments, where resolution could differ, more subject to misinterpretation. For this reason, Raw Integrated Intensity should not be used to quantify image spots.

Integrated Intensity

Integrated intensity is defined as

$$\text{Integrated Intensity} = a(\Sigma I_i - \textit{Pixels}\ \bar{b})$$

where area per pixel, a = (resolution $\times 10^{-3})^2$, and ***Pixels*** is the number of pixels enclosed by the feature. Units are counts-mm^2 (integrated pixel volume).

When background is uniform, integrated intensity is independent of both the size of feature drawn on the image and the resolution. Integrated intensity is proportional to the amount of dye-labeled antibodies on the membrane and therefore can be accurately used for quantification.

Average Intensity

When background is zero, average intensity is raw integrated intensity divided by the pixel count.

$$\text{Average Intensity} = \frac{\text{Raw Integrated Intensity}}{\textit{Pixel Count}}$$

Since average intensity is dependent on feature area, it should not be used to quantify image spots. Average intensity can be used only when an image spot has uniform intensity, for example when estimating background (with background correction turned off). Average intensity should not be used when there are both spot and background intensities within the defined feature.

Trimmed Mean

Trimmed Mean is similar to Average Intensity, except that five percent of pixels with the highest and lowest pixel intensities are excluded from the intensity summation and pixel count.

Peak Intensity

Peak intensity is the highest pixel intensity within a feature. The peak intensity value does not have background intensity subtracted.

Minimum Intensity

Minimum intensity is the lowest pixel intensity within a feature. The minimum intensity value does not have background intensity subtracted.

Signal-to-Noise Ratio

Signal-to-noise ratio is defined as follows:

$$\text{SN Ratio} = \frac{\text{Peak Intensity - Background } (\bar{b})}{\textit{Std. Dev. Background } (\bar{b})}$$

Concentration

Concentration is the amount of fluorescent material present within a given feature. Concentration is calculated relative to user-defined concentration standards on the same image. The units of concentration are always the same as the standards. To calculate concentration, the intensity of each concentration standard is plotted and fitted with a curve using one of four interpolation methods: linear, log, reciprocal fit, or exponential. Next, the concentration of unknown image spots are calculated by comparing the intensity of the area within the surrounding feature to the curve. Additional information on choosing the interpolation method is given in Chapter 8.

Probability

Probability is one of the values that can be output in a report. Probability represents the probability that the area enclosed by the feature boundary contains a statistically significant area of higher signal. Odyssey software uses a "Student's" t-Test to determine whether the mean of a set of pixels inside a given feature is significantly different from the mean of another data set of background pixels. The t-Test is defined as follows:

Students t-Test:

$$tTest = \frac{\sqrt{(n_s \times n_b \times dof)/(n_s + n_b)(a_s - a_b)}}{\sqrt{(\nu_s)(n_s - 1) + (\nu_b)(n_b - 1)}}$$

Where:

n_s	is the number of pixels in the signal area
n_b	is the number of pixels in the background area
$dof = (n_b+n_s-2)$	is the degrees of freedom
a_s	is the average of signal pixels
a_b	is the average of background pixels
v_s	is the variance of signal pixals
v_b	is the variance of background pixels

The value of the t-Test can be positive or negative. If the t-Test is negative, then the mean of pixels within the feature is less than the mean of background pixels. Since this is meaningless for Odyssey applications, Odyssey software reports "n/a" if the t-Test is negative. (Odyssey software versions prior to 1.1, however, did report negative values.). The reported probability (usually "n/a", -0.99%, or 10000%) will also be meaningless if there is no background defined or if "Lane" is selected as the background method.

When the value of the t-Test is positive and not zero, it is further refined using a published, iterative statistical algorithm that calculates a probability in percent. This algorithm was translated from a perl translation of the Pascal function on p. 81 of "Statistical Computing in Pascal" by D. Cooke, A.H. Craven, and G.M. Clark (1985: Edward Arnold (Pubs) Ltd: London). The Pascal algorithm is itself a translation of the Fortran algorithm "AS 3" by B.E. Cooper of the Atlas Computer Laboratory, as reported in (among other places) "Applied Statistical Algorithms" edited by P. Griffiths and I.D. Hill (1985: Ellis Horwood Ltd.; W. Sussex, England).

Molecular Weight

The molecular weights of bands in a lane are determined by comparison to the positions of molecular weight standard bands in lanes containing standards. Chapter 6 discusses the use of molecular

weight standards to calibrate the molecular weight of unknown bands.

Percent Saturation

Percent saturation is one of the values that can be output on reports (Chapter 10). Percent saturation is defined as the number of saturated pixels divided by the total number of pixels enclosed in a feature, multiplied by 100 to give a percentage.

Percent Response for ICW Assays

When ICW calculations start, all wells designated as background (see Chapter 9) are averaged for the 700-channel image. Similarly, the 800-channel background wells are averaged, resulting in each channel having its own background intensity value. Assuming that background subtraction is enabled, which it normally should be, the average background intensity for the 700 channel is subtracted from the integrated intensity of each well in the 700 channel. The same calculation is performed on the wells of the 800-channel image using its corresponding background and integrated intensity values. References to integrated intensity throughout the rest of this discussion refer to the original integrated intensity minus background intensity.

Odyssey software allows complete flexibility in how the image channels are used, but suppose the 700 channel is used to detect phosphorylated proteins and the 800 channel is used to detect total protein. In this example, the 800 channel would be designated as the channel used to calculate relative intensity, which indicates the relative number of cells. In the channel used to calculate relative intensity, Odyssey software starts by finding the well with the maximum integrated intensity. All wells designated as **Sample** or **100% Standard** in the 800 channel are divided by the maximum

integrated intensity to obtain the relative intensity of each well. The relative intensity values will normally be between 0.0 and 1.0, though negative numbers are possible. (Negative relative intensities indicate the original integrated intensity was lower than the average background when the background was subtracted). Relative intensity values with low statistical significance are color coded when displayed in the View ICW Analysis window.

Continuing the example, the next step is to use the relative intensity values from the 800 channel to normalize integrated intensity values in the 700 channel. To normalize the 700 channel, the integrated intensity for each well in the 700 channel is divided by the relative intensity value from the corresponding well in the 800 channel. Next, all wells designated **100% Standard** in the 700 channel are averaged. The normalized value for each well is then divided by the **100% Standard** of the 700 channel and multiplied by 100 to give a value that is the percentage response to the control in the **100% Standard**.

The calculation is slightly different if rows are linked. When rows are linked, all the integrated intensity values for the linked wells in a given column are averaged. The average integrated intensity replaces the original integrated intensity in each of the linked wells.

Note: *If the* ***Calculate Relative Intensity in Channel*** *field in the ICW parameters is deselected (Chapter 9), the percent response will be calculated for both channels with no normalization. For each channel, the relative intensity values are divided by the 100% Standard and multiplied by 100.*

ICW calculations are performed as described above when a whole microplate is analyzed. If the **Individual Row Analyses** or I**ndividual Column Analyses** check boxes are checked on the Calculations tab of the ICW Parameters window, **%Response** is calculated using only the wells within a given row or column. For example, when **Individual Row Analyses** is checked, each row analyzed must have at least one **Background** well and **100% Standard** well. **%Response** for sample wells in the row is calculated using the **Background** and

100% Standard well(s) in the same row as the sample wells. An error message will be displayed if **Background** and **100% Standard** wells have not been designated. If there is more than one **Background** or **100% Standard** well in a row, they are averaged before being used in the calculations, but they are not averaged with any **Background** or **100% Standard** wells outside the row. When individual rows are analyzed, wells between rows cannot be linked, so any links designated on the **Well Links** tab are ignored. When analyzing individual columns, the same concept applies: **Background** and **100% Standard** wells must be designated in each column to be analyzed and columns cannot be linked.

Z'-Factor Calculations

Before an assay can be used with various test compounds, the assay should be optimized for reagents, protocols, instrumentation, kinetics, and other conditions not related to the test compounds. Calculating the Z'-factor* for an assay provides a method to evaluate whether assay conditions are optimized. Z'-factor is a dimensionless value that indicates whether the assay has sufficient dynamic range and low enough data variability to generate meaningful data.

When Z'-factor calculations start, all wells designated as background in the 700 channel are averaged and the same is done for the 800 channel. Assuming that background subtraction is enabled, which it normally should be, the average background intensity for the 700 channel is subtracted from the integrated intensity of all 700-channel wells. The background is also subtracted from the 800 channel image.

During setup, one channel is designated as the channel used to calculate relative intensity. This channel is used to normalize for sample loading by detecting a housekeeping protein or protein stain. In the channel used to calculate relative intensity, Odyssey software starts by finding the well with the maximum integrated intensity. All

wells in this channel are divided by the maximum integrated intensity to obtain the relative intensity of each well. The relative intensity values will normally be between 0.0 and 1.0, though negative numbers are possible. Relative intensity values with low statistical significance are color coded when displayed in the View ICW Analysis window.

The next step is to use the relative intensity values to normalize integrated intensity values in the sample channel. The integrated intensity for each well in the sample channel is divided by the relative intensity value from the corresponding well in the channel used to calculate relative intensity.

The wells designated positive and negative controls are now ready for the Z'-factor calculations. Z'-factor is calculated according to the following equation:

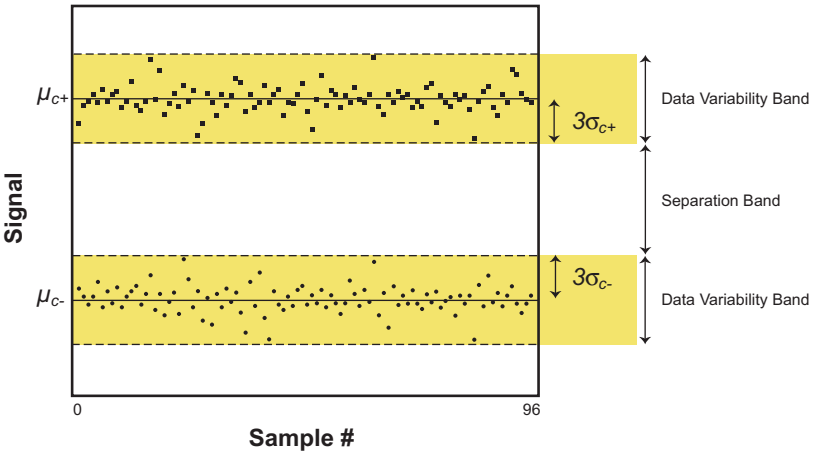
$$Z' = \frac{|\mu_{c+} - \mu_{c-}| - (3\sigma_{c+} + 3\sigma_{c-})}{|\mu_{c+} - \mu_{c-}|} = 1 - \frac{(3\sigma_{c+} + 3\sigma_{c-})}{|\mu_{c+} - \mu_{c-}|}$$

Where:
$3\sigma_{c+}$ = 3 standard deviations of positive controls
$3\sigma_{c-}$ = 3 standard deviations of negative controls
μ_{c+} is the mean of the positive controls (maximum signal)
μ_{c-} is the mean of the negative controls (minimum signal)

and

$|\mu_{c+} - \mu_{c-}|$ is the assay dynamic range
$|\mu_{c+} - \mu_{c-}| - (3\sigma_{c+} + 3\sigma_{c-})$ is the separation band between positive and negative control signals.

These variables are represented in the illustration below.



Dimensionless Z'-factor values are always 1 or less as described below.

- Z' = 1 is an ideal assay. As standard deviations become very small or the difference between signals for positive and negative controls approaches infinity, Z'-factor approaches 1.
- 1 > Z' ≥ 0.5 indicates a high quality assay exhibiting a wide separation between signals for positive and negative controls, and low data variability.
- 0.5 >Z' > 0 may indicate a low quality assay with marginal distinction between signals for positive and negative controls, and higher data variability. However, an acceptable Z'-factor target value should be determined prior to performing final validation of an assay. An assay with a relatively low number of data points, such as the number obtained from a 96-well plate, may produce a Z'-factor value less than 0.5 but still be considered a good quality assay if the same value is acheived between different plates run on different days.
- Z' ≤ 0 indicates unreliable data.

A Z'-factor value that is close to zero or negative indicates that assay conditions are not optimized or the assay is not capable of generating meaningful data.

* Zhang, J.-H., Chung, T.D., and Oldenburg, K.R. 1999. A Simple Statistical Parameter for Use in Evaluation and Validation of High Throughput Screening Assays. J. Biomol. Screen. 4: 2, 67-73.

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